



SEQUENCE LISTING

Schuetz, John <120> A Method of Identifying and/or Isolating Stem
Cells <130> 1340-1-021CIP2 <140> 09/866,866 <141> 2001-05-29 <150>
09/584,586 <151> 2000-05-31 <150> PCT/US99/11825 <151> 1999-05-27
<150> 60/086,988 <151> 1998-05-28 <160> 27 <170> PatentIn
version 3.0 <210> 1 <211> 3860 <212> DNA <213> Homo sapiens <400>

1
atggatcttg aaggggaccg caatggagga gcaaagaaga agaacttttt taaactgaac 60
aataaaagtg aaaaagataa gaaggaaaag aaaccaactg tcagtgtatt ttcaatgttt 120
cgctattcaa attggcttga caagttgtat atgggtggtg gaactttggc tgccatcatc 180
catggggctg gacttcctct catgatgctg gtgtttggag aaatgacaga tatctttgca 240
aatgcaggaa atttagaaga tctgatgtca aacatcacta atagaagtga tatcaatgat 300
acagggttct tcatgaatct ggaggaagac atgaccagat atgcctatta ttacagtgga 360
attggtgctg gggtgctggt tgctgcttac attcaggttt cattttggtg cctggcagct 420
ggaagacaaa tacacaaaat tagaaaacag ttttttcatg ctataatgcg acaggagata 480
ggctggtttg atgtgcacga tgttggggag cttaacaccc gacttacaga tgatgtctct 540
aagattaatg aaggatattg tgacaaaatt ggaatgttct ttcagtcaat ggcaacattt 600
ttcactgggt ttatagtagg atttacacgt ggttggaagc taacccttgt gattttggcc 660
atcagtctcg ttcttggact gtcagctgct gtctgggcaa agatactatc ttcatttact 720
gataaagaac tcttagcgta tgcaaaagct ggagcagtag ctgaagaggt cttggcagca 780
attagaactg tgattgcatt tggaggacaa aagaaagaac ttgaaaggta caacaaaaat 840
ttagaagaag ctaaaagaat tgggataaag aaagctatta cagccaatat ttctataggt 900
gctgctttcc tgctgatcta tgcactttat gctctggcct tctggtatgg gaccaccttg 960
gtcctctcag gggaatattc tattggacaa gtactcactg tattcttttc tgtattaatt 1020
ggggctttta gtgttggaac ggcactctca agcattgaag catttgcaaa tgcaagagga 1080
gcagcttatg aaatcttcaa gataattgat aataagccaa gtattgacag ctattcgaag 1140
agtgggcaca aaccagataa tattaaggga aatttggaat tcagaaatgt tcaattcagt 1200
taccatctc gaaaagaagt taagatcttg aagggcctga acctgaaggt gcagagtggg 1260
cagacgggtg ccctgggttg aaacagtggc tgtgggaaga gcacaacagt ccagctgatg 1320
cagaggctct atgacccac agaggggatg gtcagtgttg atggacagga tattaggacc 1380
ataaatgtaa ggtttctacg ggaaatcatt ggtgtggtga gtcaggaacc tgtattgttt 1440
gccaccacga tagctgaaaa cattcgctat ggccgtgaaa atgtcaccat ggatgagatt 1500
gagaaagctg tcaaggaagc caatgcctat gactttatca tgaaactgcc tcataaattt 1560
gacaccctgg ttggagagag aggggccag ttgagtgtg ggcagaagca gaggatcgcc 1620
attgcacgtg ccctggttcg caaccccaag atcctcctgc tggatgaggc cacgtcagcc 1680
ttggacacag aaagcgaagc agtggttcag gtggctctgg ataaggccag aaaaggctcg 1740
accaccattg tgatagctca tcgtttgtct acagttcgta atgctgacgt catcgctggt 1800

ttcgatgatg	gagtcattgt	ggagaaagga	aatcatgatg	aactcatgaa	agagaaaggc	1860
atttacttca	aacttgtcac	aatgcagaca	gcaggaaatg	aagttgaatt	agaaaatgca	1920
gctgatgaat	ccaaaagtga	aattgatgcc	ttggaaatgt	cttcaaatga	ttcaagatcc	1980
agtctaataa	gaaaaagatc	aactcgtagg	agtgtccgtg	gatcacaagc	ccaagacaga	2040
aagcttagta	ccaaagaggc	tctggatgaa	agtatacctc	cagtttcctt	ttggaggatt	2100
atgaagctaa	atttaactga	atggccttat	tttgttgttg	gtgtattttg	tgccattata	2160
aatggaggcc	tgcaaccagc	atttgcaata	atattttcaa	agattatagg	ggtttttaca	2220
agaattgatg	atcctgaaac	aaaacgacag	aatagtaact	tgttttcact	attgttttcta	2280
gcccttggaa	ttattttctt	tattacattt	ttccttcaag	gtttcacatt	tggcaaagct	2340
ggagagatcc	tcaccaagcg	gctccgatac	atggttttcc	gatccatgct	cagacaggat	2400
gtgagttggt	ttgatgaccc	taaaaacacc	actggagcat	tgactaccag	gctcgccaat	2460
gatgctgctc	aagttaaagg	ggctataggt	tccaggcttg	ctgtaattac	ccagaatata	2520
gcaaactctt	ggacaggaat	aattatatcc	ttcatctatg	gttggcaact	aacactgtta	2580
ctcttagcaa	ttgtacccat	cattgcaata	gcaggagttg	ttgaaatgaa	aatgtttgtct	2640
ggacaagcac	tgaaagataa	gaaagaacta	gaaggtgctg	ggaagatcgc	tactgaagca	2700
atagaaaact	tccgaaccgt	tgtttctttg	actcaggagc	agaagtttga	acatatgtat	2760
gctcagagtt	tgccaggtacc	atacagaaac	tctttgagga	aagcacacat	ctttggaatt	2820
acattttcct	tcaccagggc	aatgatgtat	ttttcctatg	ctggatgttt	ccggtttgga	2880
gcctacttgg	tggcacataa	actcatgagc	tttgaggatg	ttctgttagt	attttcagct	2940
gttgtctttg	gtgccatggc	cgtggggcaa	gtcagttcat	ttgctcctga	ctatgccaaa	3000
gccaaaatat	cagcagccca	catcatcatg	atcattgaaa	aaaccctttt	gattgacagc	3060
tacagcacgg	aaggccta	gccgaacaca	ttggaaggaa	atgtcacatt	tggatgaagtt	3120
gtattcaact	atcccacccg	accggacatc	ccagtgtctc	agggactgag	cctggagggtg	3180
aagaagggcc	agacgctggc	tctggtgggc	agcagtggct	gtgggaagag	cacagtggtc	3240
cagctcctgg	agcggttcta	cgacccttg	gcagggaag	tgctgcttga	tggcaaagaa	3300
ataaagcgac	tgaatgttca	gtggctccga	gcacacctgg	gcacgtgtgc	ccaggagccc	3360
atcctgtttg	actgcagcat	tgctgagaac	attgcctatg	gagacaacag	ccgggtggtg	3420
tcacaggaag	agatcgtgag	ggcagcaaag	gaggccaaca	tacatgcctt	catcgagtca	3480
ctgcctaata	aatatagcac	taaagtagga	gacaaaggaa	ctcagctctc	tgggtggccag	3540
aaacaacgca	ttgccatagc	tcgtgccctt	gtagacagc	ctcatatttt	gcttttggat	3600
gaagccacgt	cagctctgga	tacagaaagt	gaaaagggtg	tccaagaagc	cctggacaaa	3660
gccagagaag	gccgcacctg	cattgtgatt	gctcacccgc	tgtccaccat	ccagaatgca	3720
gacttaatag	tggtgtttca	gaatggcaga	gtcaaggagc	atggcacgca	tcagcagctg	3780

ctggcacaga aaggcatcta tttttcaatg gtcagtgtcc aggctggaac aaagcgccag 3840
tgaactctgg ttaactccac 3860

<210>	2	<211>	1280	<212>	PRT	<213>	Homo sapiens	<400>	2						
Met 1	Asp	Leu	Glu	Gly 5	Asp	Arg	Asn	Gly	Gly 10	Ala	Lys	Lys	Lys	Asn 15	Phe
Phe	Lys	Leu	Asn 20	Asn	Lys	Ser	Glu	Lys 25	Asp	Lys	Lys	Glu	Lys 30	Lys	Pro
Thr	Val	Ser 35	Val	Phe	Ser	Met	Phe 40	Arg	Tyr	Ser	Asn	Trp 45	Leu	Asp	Lys
Leu	Tyr 50	Met	Val	Val	Gly	Thr 55	Leu	Ala	Ala	Ile	Ile 60	His	Gly	Ala	Gly
Leu 65	Pro	Leu	Met	Met	Leu	Val	Phe	Gly	Glu	Met 75	Thr	Asp	Ile	Phe	Ala 80
Asn	Ala	Gly	Asn 85	Leu	Glu	Asp	Leu	Met	Ser 90	Asn	Ile	Thr	Asn	Arg 95	Ser
Asp	Ile	Asn	Asp 100	Thr	Gly	Phe	Phe	Met 105	Asn	Leu	Glu	Glu	Asp	Met	Thr
Arg	Tyr	Ala 115	Tyr	Tyr	Tyr	Ser	Gly 120	Ile	Gly	Ala	Gly	Val 125	Leu	Val	Ala
Ala	Tyr 130	Ile	Gln	Val	Ser	Phe 135	Trp	Cys	Leu	Ala	Ala 140	Gly	Arg	Gln	Ile
His 145	Lys	Ile	Arg	Lys	Gln 150	Phe	Phe	His	Ala	Ile 155	Met	Arg	Gln	Glu	Ile 160
Gly	Trp	Phe	Asp	Val 165	His	Asp	Val	Gly	Glu 170	Leu	Asn	Thr	Arg	Leu 175	Thr
Asp	Asp	Val	Ser 180	Lys	Ile	Asn	Glu	Gly 185	Ile	Gly	Asp	Lys	Ile 190	Gly	Met
Phe	Phe	Gln 195	Ser	Met	Ala	Thr	Phe 200	Phe	Thr	Gly	Phe	Ile 205	Val	Gly	Phe
Thr	Arg 210	Gly	Trp	Lys	Leu	Thr 215	Leu	Val	Ile	Leu	Ala 220	Ile	Ser	Pro	Val
Leu 225	Gly	Leu	Ser	Ala	Ala 230	Val	Trp	Ala	Lys	Ile 235	Leu	Ser	Ser	Phe	Thr 240
Asp	Lys	Glu	Leu	Leu 245	Ala	Tyr	Ala	Lys	Ala 250	Gly	Ala	Val	Ala	Glu 255	Glu
Val	Leu	Ala	Ala 260	Ile	Arg	Thr	Val	Ile 265	Ala	Phe	Gly	Gly	Gln 270	Lys	Lys
Glu	Leu	Glu 275	Arg	Tyr	Asn	Lys	Asn 280	Leu	Glu	Glu	Ala	Lys 285	Arg	Ile	Gly
Ile	Lys 290	Lys	Ala	Ile	Thr	Ala 295	Asn	Ile	Ser	Ile	Gly 300	Ala	Ala	Phe	Leu
Leu	Ile	Tyr	Ala	Ser	Tyr	Ala	Leu	Ala	Phe	Trp	Tyr	Gly	Thr	Thr	Leu

305		310		315		320
Val Leu Ser Gly	Glu Tyr Ser Ile Gly	Gln Val Leu Thr Val	Phe Phe			
	325	330	335			
Ser Val Leu Ile	Gly Ala Phe Ser Val	Gly Gln Ala Ser	Pro Ser Ile			
	340	345	350			
Glu Ala Phe Ala	Asn Ala Arg Gly	Ala Ala Tyr Glu	Ile Phe Lys Ile			
	355	360	365			
Ile Asp Asn Lys	Pro Ser Ile Asp	Ser Tyr Ser Lys	Ser Gly His Lys			
	370	375	380			
Pro Asp Asn Ile	Lys Gly Asn Leu	Glu Phe Arg Asn	Val His Phe Ser			
	385	390	395			400
Tyr Pro Ser Arg	Lys Glu Val Lys	Ile Leu Lys Gly	Leu Asn Leu Lys			
	405	410	415			
Val Gln Ser Gly	Gln Thr Val Ala	Leu Val Gly Asn	Ser Gly Cys Gly			
	420	425	430			
Lys Ser Thr Thr	Val Gln Leu Met	Gln Arg Leu Tyr	Asp Pro Thr Glu			
	435	440	445			
Gly Met Val Ser	Val Asp Gly Gln	Asp Ile Arg Thr	Ile Asn Val Arg			
	450	455	460			
Phe Leu Arg Glu	Ile Ile Gly Val	Val Ser Gln Glu	Pro Val Leu Phe			
	465	470	475			480
Ala Thr Thr Ile	Ala Glu Asn Ile	Arg Tyr Gly Arg	Glu Asn Val Thr			
	485	490	495			
Met Asp Glu Ile	Glu Lys Ala Val	Lys Glu Ala Asn	Ala Tyr Asp Phe			
	500	505	510			
Ile Met Lys Leu	Pro His Lys Phe	Asp Thr Leu Val	Gly Glu Arg Gly			
	515	520	525			
Ala Gln Leu Ser	Gly Gly Gln Lys	Gln Arg Ile Ala	Ile Ala Arg Ala			
	530	535	540			
Leu Val Arg Asn	Pro Lys Ile Leu	Leu Leu Asp Glu	Ala Thr Ser Ala			
	545	550	555			560
Leu Asp Thr Glu	Ser Glu Ala Val	Val Gln Val Ala	Leu Asp Lys Ala			
	565	570	575			
Arg Lys Gly Arg	Thr Thr Ile Val	Ile Ala His Arg	Leu Ser Thr Val			
	580	585	590			
Arg Asn Ala Asp	Val Ile Ala Gly	Phe Asp Asp Gly	Val Ile Val Glu			
	595	600	605			
Lys Gly Asn His	Asp Glu Leu Met	Lys Glu Lys Gly	Ile Tyr Phe Lys			
	610	615	620			
Leu Val Thr Met	Gln Thr Ala Gly	Asn Glu Val Glu	Leu Glu Asn Ala			
	625	630	635			640
Ala Asp Glu Ser	Lys Ser Glu Ile	Asp Ala Leu Glu	Met Ser Ser Asn			
	645	650	655			
Asp Ser Arg Ser	Ser Leu Ile Arg	Lys Arg Ser Thr	Arg Arg Ser Val			

660						665						670					
Arg	Gly	Ser	Gln	Ala	Gln	Asp	Arg	Lys	Leu	Ser	Thr	Lys	Glu	Ala	Leu		
Asp	Glu	Ser	Ile	Pro	Pro	Val	Ser	Phe	Trp	Arg	Ile	Met	Lys	Leu	Asn		
Leu	Thr	Glu	Trp	Pro	Tyr	Phe	Val	Val	Gly	Val	Phe	Cys	Ala	Ile	Ile		
Asn	Gly	Gly	Leu	Gln	Pro	Ala	Phe	Ala	Ile	Ile	Phe	Ser	Lys	Ile	Ile		
Gly	Val	Phe	Thr	Arg	Ile	Asp	Asp	Pro	Glu	Thr	Lys	Arg	Gln	Asn	Ser		
Asn	Leu	Phe	Ser	Leu	Leu	Phe	Leu	Ala	Leu	Gly	Ile	Ile	Ser	Phe	Ile		
Thr	Phe	Phe	Leu	Gln	Gly	Phe	Thr	Phe	Gly	Lys	Ala	Gly	Glu	Ile	Leu		
Thr	Lys	Arg	Leu	Arg	Tyr	Met	Val	Phe	Arg	Ser	Met	Leu	Arg	Gln	Asp		
Val	Ser	Trp	Phe	Asp	Asp	Pro	Lys	Asn	Thr	Thr	Gly	Ala	Leu	Thr	Thr		
Arg	Leu	Ala	Asn	Asp	Ala	Ala	Gln	Val	Lys	Gly	Ala	Ile	Gly	Ser	Arg		
Leu	Ala	Val	Ile	Thr	Gln	Asn	Ile	Ala	Asn	Leu	Gly	Thr	Gly	Ile	Ile		
Ile	Ser	Phe	Ile	Tyr	Gly	Trp	Gln	Leu	Thr	Leu	Leu	Leu	Leu	Ala	Ile		
Val	Pro	Ile	Ile	Ala	Ile	Ala	Gly	Val	Val	Glu	Met	Lys	Met	Leu	Ser		
Gly	Gln	Ala	Leu	Lys	Asp	Lys	Lys	Glu	Leu	Glu	Gly	Ala	Gly	Lys	Ile		
Ala	Thr	Glu	Ala	Ile	Glu	Asn	Phe	Arg	Thr	Val	Val	Ser	Leu	Thr	Gln		
Glu	Gln	Lys	Phe	Glu	His	Met	Tyr	Ala	Gln	Ser	Leu	Gln	Val	Pro	Tyr		
Arg	Asn	Ser	Leu	Arg	Lys	Ala	His	Ile	Phe	Gly	Ile	Thr	Phe	Ser	Phe		
Thr	Gln	Ala	Met	Met	Tyr	Phe	Ser	Tyr	Ala	Gly	Cys	Phe	Arg	Phe	Gly		
Ala	Tyr	Leu	Val	Ala	His	Lys	Leu	Met	Ser	Phe	Glu	Asp	Val	Leu	Leu		
Val	Phe	Ser	Ala	Val	Val	Phe	Gly	Ala	Met	Ala	Val	Gly	Gln	Val	Ser		
Ser	Phe	Ala	Pro	Asp	Tyr	Ala	Lys	Ala	Lys	Ile	Ser	Ala	Ala	His	Ile		
Ile	Met	Ile	Ile	Glu	Lys	Thr	Pro	Leu	Ile	Asp	Ser	Tyr	Ser	Thr			

1010	1015	1020
Glu Gly Leu Met Pro Asn Thr 1025	Leu Glu Gly Asn Val 1030	Thr Phe Gly 1035
Glu Val Val Phe Asn Tyr Pro 1040	Thr Arg Pro Asp Ile 1045	Pro Val Leu 1050
Gln Gly Leu Ser Leu Glu Val 1055	Lys Lys Gly Gln Thr 1060	Leu Ala Leu 1065
Val Gly Ser Ser Gly Cys Gly 1070	Lys Ser Thr Val Val 1075	Gln Leu Leu 1080
Glu Arg Phe Tyr Asp Pro Leu 1085	Ala Gly Lys Val Leu 1090	Leu Asp Gly 1095
Lys Glu Ile Lys Arg Leu Asn 1100	Val Gln Trp Leu Arg 1105	Ala His Leu 1110
Gly Ile Val Ser Gln Glu Pro 1115	Ile Leu Phe Asp Cys 1120	Ser Ile Ala 1125
Glu Asn Ile Ala Tyr Gly Asp 1130	Asn Ser Arg Val Val 1135	Ser Gln Glu 1140
Glu Ile Val Arg Ala Ala Lys 1145	Glu Ala Asn Ile His 1150	Ala Phe Ile 1155
Glu Ser Leu Pro Asn Lys Tyr 1160	Ser Thr Lys Val Gly 1165	Asp Lys Gly 1170
Thr Gln Leu Ser Gly Gly Gln 1175	Lys Gln Arg Ile Ala 1180	Ile Ala Arg 1185
Ala Leu Val Arg Gln Pro His 1190	Ile Leu Leu Leu Asp 1195	Glu Ala Thr 1200
Ser Ala Leu Asp Thr Glu Ser 1205	Glu Lys Val Val Gln 1210	Glu Ala Leu 1215
Asp Lys Ala Arg Glu Gly Arg 1220	Thr Cys Ile Val Ile 1225	Ala His Arg 1230
Leu Ser Thr Ile Gln Asn Ala 1235	Asp Leu Ile Val Val 1240	Phe Gln Asn 1245
Gly Arg Val Lys Glu His Gly 1250	Thr His Gln Gln Leu 1255	Leu Ala Gln 1260
Lys Gly Ile Tyr Phe Ser Met 1265	Val Ser Val Gln Ala 1270	Gly Thr Lys 1275
Arg Gln 1280		

```

<210> 3 <211> 3860 <212> DNA <213> homo sapiens <400> 3
atggatcttg aaggggaccg caatggagga gcaaagaaga agaacttttt taaactgaac      60
aataaaagtg aaaaagataa gaaggaaaag aaaccaactg tcagtgtatt ttcaatgttt      120
cgctattcaa attggcttga caagttgtat atggtggtgg gaactttggc tgccatcatc      180
catggggctg gacttcctct catgatgctg gtgtttggag aaatgacaga tatctttgca      240
aatgcaggaa atttagaaga tctgatgtca aacatcacta atagaagtga tatcaatgat      300

```

acagggttct	tcatgaatct	ggaggaagac	atgaccagat	atgcctatta	ttacagtgga	360
attgggtgctg	gggtgctggt	tgctgcttac	attcaggttt	catttttggtg	cctggcagct	420
ggaagacaaa	tacacaaaat	tagaaaacag	ttttttcatg	ctataatgcg	acaggagata	480
ggctggtttg	atgtgcacga	tggtggggag	cttaacaccc	gacttacaga	tgatgtctct	540
aagattaatg	aagttatttg	tgacaaaatt	ggaatgttct	ttcagtcaat	ggcaacattt	600
ttcactgggt	ttatagtagg	atttacacgt	ggttggaagc	taacccttgt	gattttggcc	660
atcagtctctg	ttcttggaact	gtcagctgct	gtctgggcaa	agatactatc	ttcatttact	720
gataaagaac	tcttagcgta	tgcaaaagct	ggagcagtag	ctgaagaggt	cttggcagca	780
attagaactg	tgattgcatt	tggaggacaa	aagaaagaac	ttgaaaggta	caacaaaaat	840
ttagaagaag	ctaaaagaat	tgggataaag	aaagctatta	cagccaatat	ttctataggt	900
gctgctttcc	tgctgatcta	tgcatcttat	gctctggcct	tctggtatgg	gaccaccttg	960
gtcctctcag	gggaatatct	tattggacaa	gtactcactg	tattcttttc	tgtattaatt	1020
ggggctttta	gtgttggaac	ggcatctcca	agcattgaag	catttgcaaa	tgcaagagga	1080
gcagcttatg	aaatcttcaa	gataattgat	aataagccaa	gtattgacag	ctattcgaag	1140
agtgggcaca	aaccagataa	tattaaggga	aatttggaat	tcagaaatgt	tcacttcagt	1200
tacccatctc	gaaaagaagt	taagatcttg	aagggcctga	acctgaaggt	gcagagtggg	1260
cagacgggtg	ccctggttg	aaacagtggc	tgtgggaaga	gcacaacagt	ccagctgatg	1320
cagaggctct	atgaccccac	agaggggatg	gtcagtgttg	atggacagga	tattaggacc	1380
ataaatgtaa	ggtttctacg	ggaaatcatt	gggtgtggtga	gtcaggaacc	tgtattgttt	1440
gccaccacga	tagctgaaaa	cattcgctat	ggccgtgaaa	atgtcaccat	ggatgagatt	1500
gagaaagctg	tcaaggaagc	caatgcctat	gactttatca	tgaaactgcc	tcataaatct	1560
gacaccctgg	ttggagagag	aggggccag	ttgagtgggtg	ggcagaagca	gaggatcgcc	1620
attgcacgtg	ccctggttcg	caaccccaag	atcctcctgc	tggatgaggc	cacgtcagcc	1680
ttggacacag	aaagcgaagc	agtggttcag	gtggctcttg	ataaggccag	aaaaggtcgg	1740
accaccattg	tgatagctca	tcgtttgtct	acagttcgta	atgctgacgt	catcgctggt	1800
ttcgatgatg	gagtcattgt	ggagaaagga	aatcatgatg	aactcatgaa	agagaaaggc	1860
atttacttca	aacttgtcac	aatgcagaca	gcaggaaatg	aagttgaatt	agaaaatgca	1920
gctgatgaat	ccaaaagtga	aattgatgcc	ttggaaatgt	cttcaaatga	ttcaagatcc	1980
agtctaataa	gaaaaagatc	aactcgtagg	agtgtccgtg	gatcacaagc	ccaagacaga	2040
aagcttagta	ccaaagaggc	tctggatgaa	agtatacctc	cagtttcctt	ttggaggatt	2100
atgaagctaa	atttaactga	atggccttat	ttgtttgttg	gtgtattttg	tgccattata	2160
aatggaggcc	tgcaaccagc	atttgcaata	atattttcaa	agattatagg	ggtttttaca	2220
agaattgatg	atcctgaaac	aaaacgacag	aatagtaact	tgttttcact	attgtttcta	2280

```

gcccttggaa ttatttcttt tattacattt ttccttcaag gtttcacatt tggcaaagct 2340
ggagagatcc tcaccaagcg gtcctgatac atgggttttcc gatccatgct cagacaggat 2400
gtgagttggt ttgatgaccc taaaaacacc actggagcat tgactaccag gctcgccaat 2460
gatgctgctc aagttaaagg ggctataggt tccaggcttg ctgtaattac ccagaatata 2520
gcaaattcttg ggacaggaat aattatatcc ttcattctatg gttggcaact aacactgtta 2580
ctcttagcaa ttgtacccat cattgcaata gcaggagttg ttgaaatgaa aatgttgtct 2640
ggacaagcac tgaaagataa gaaagaacta gaagggtgctg ggaagatcgc tactgaagca 2700
atagaaaact tccgaaccgt tgtttctttg actcaggagc agaagtttga acatatgtat 2760
gctcagagtt tgcagggtacc atacagaaac tctttgagga aagcacacat ctttggaatt 2820
acattttcct tcacccaggc aatgatgtat ttttcctatg ctggatgttt ccggtttgga 2880
gcctacttgg tggcacataa actcatgagc tttgaggatg ttctgttagt attttcagct 2940
gttgtctttg gtgccatggc cgtggggcaa gtcagttcat ttgctcctga ctatgccaaa 3000
gccaaaatat cagcagccca catcatcatg atcattgaaa aaacccttt gattgacagc 3060
tacagcacgg aaggcctaata gccgaacaca ttggaaggaa atgtcacatt tgggtgaagtt 3120
gtattcaact atcccacccg accggacatc ccagtgttc agggactgag cctggaggtg 3180
aagaagggcc agacgctggc tctggtgggc agcagtggct gtgggaagag cacagtggtc 3240
cagctcctgg agcggttcta cgacccttg gcagggaaag tgctgcttga tggcaaagaa 3300
ataaagcgac tgaatgttca gtggctccga gcacacctgg gcacgtgtc ccaggagccc 3360
atcctgtttg actgcagcat tgctgagaac attgcctatg gagacaacag ccgggtggtg 3420
tcacaggaag agatcgtgag ggcagcaaag gaggccaaca tacatgcctt catcgagtca 3480
ctgcctaata aatatagcac taaagtagga gacaaaggaa ctcagctctc tgggtggccag 3540
aaacaacgca ttgccatagc tcgtgccctt gttagacagc ctcatatttt gcttttggat 3600
gaagccacgt cagctctgga tacagaaagt gaaaagggtt tccaagaagc cctggacaaa 3660
gccagagaag gccgcacctg cattgtgatt gtcaccgcc tgtccaccat ccagaatgca 3720
gacttaatag tgggtgttca gaatggcaga gtcaaggagc atggcacgca tcagcagctg 3780
ctggcacaga aaggcatcta tttttcaatg gtcagtgtcc aggctggaac aaagcgccag 3840
tgaactctgg ttaactccac 3860

```

<210> 4 <211> 1280 <212> PRT <213> Homo sapiens <400> 4

Met Asp Leu Glu Gly Asp Arg Asn Gly Gly Ala Lys Lys Lys Asn Phe
1 5 10 15

Phe Lys Leu Asn Asn Lys Ser Glu Lys Asp Lys Lys Glu Lys Lys Pro
20 25 30

Thr Val Ser Val Phe Ser Met Phe Arg Tyr Ser Asn Trp Leu Asp Lys
35 40 45

Leu	Tyr	Met	Val	Val	Gly	Thr	Leu	Ala	Ala	Ile	Ile	His	Gly	Ala	Gly	
50						55					60					
Leu	Pro	Leu	Met	Met	Leu	Val	Phe	Gly	Glu	Met	Thr	Asp	Ile	Phe	Ala	
65					70					75					80	
Asn	Ala	Gly	Asn	Leu	Glu	Asp	Leu	Met	Ser	Asn	Ile	Thr	Asn	Arg	Ser	
			85						90					95		
Asp	Ile	Asn	Asp	Thr	Gly	Phe	Phe	Met	Asn	Leu	Glu	Glu	Asp	Met	Thr	
			100					105					110			
Arg	Tyr	Ala	Tyr	Tyr	Tyr	Ser	Gly	Ile	Gly	Ala	Gly	Val	Leu	Val	Ala	
	115						120					125				
Ala	Tyr	Ile	Gln	Val	Ser	Phe	Trp	Cys	Leu	Ala	Ala	Gly	Arg	Gln	Ile	
	130					135					140					
His	Lys	Ile	Arg	Lys	Gln	Phe	Phe	His	Ala	Ile	Met	Arg	Gln	Glu	Ile	
145					150					155					160	
Gly	Trp	Phe	Asp	Val	His	Asp	Val	Gly	Glu	Leu	Asn	Thr	Arg	Leu	Thr	
				165					170					175		
Asp	Asp	Val	Ser	Lys	Ile	Asn	Glu	Val	Ile	Gly	Asp	Lys	Ile	Gly	Met	
			180					185					190			
Phe	Phe	Gln	Ser	Met	Ala	Thr	Phe	Phe	Thr	Gly	Phe	Ile	Val	Gly	Phe	
		195					200					205				
Thr	Arg	Gly	Trp	Lys	Leu	Thr	Leu	Val	Ile	Leu	Ala	Ile	Ser	Pro	Val	
	210					215					220					
Leu	Gly	Leu	Ser	Ala	Ala	Val	Trp	Ala	Lys	Ile	Leu	Ser	Ser	Phe	Thr	
225					230					235					240	
Asp	Lys	Glu	Leu	Leu	Ala	Tyr	Ala	Lys	Ala	Gly	Ala	Val	Ala	Glu	Glu	
				245					250					255		
Val	Leu	Ala	Ala	Ile	Arg	Thr	Val	Ile	Ala	Phe	Gly	Gly	Gln	Lys	Lys	
			260					265					270			
Glu	Leu	Glu	Arg	Tyr	Asn	Lys	Asn	Leu	Glu	Glu	Ala	Lys	Arg	Ile	Gly	
		275					280					285				
Ile	Lys	Lys	Ala	Ile	Thr	Ala	Asn	Ile	Ser	Ile	Gly	Ala	Ala	Phe	Leu	
	290					295					300					
Leu	Ile	Tyr	Ala	Ser	Tyr	Ala	Leu	Ala	Phe	Trp	Tyr	Gly	Thr	Thr	Leu	
305					310					315					320	
Val	Leu	Ser	Gly	Glu	Tyr	Ser	Ile	Gly	Gln	Val	Leu	Thr	Val	Phe	Phe	
				325					330					335		
Ser	Val	Leu	Ile	Gly	Ala	Phe	Ser	Val	Gly	Gln	Ala	Ser	Pro	Ser	Ile	
			340					345					350			
Glu	Ala	Phe	Ala	Asn	Ala	Arg	Gly	Ala	Ala	Tyr	Glu	Ile	Phe	Lys	Ile	
		355					360					365				
Ile	Asp	Asn	Lys	Pro	Ser	Ile	Asp	Ser	Tyr	Ser	Lys	Ser	Gly	His	Lys	
	370					375					380					
Pro	Asp	Asn	Ile	Lys	Gly	Asn	Leu	Glu	Phe	Arg	Asn	Val	His	Phe	Ser	
385					390					395					400	

Tyr	Pro	Ser	Arg	Lys	Glu	Val	Lys	Ile	Leu	Lys	Gly	Leu	Asn	Leu	Lys		
				405					410					415			
Val	Gln	Ser	Gly	Gln	Thr	Val	Ala	Leu	Val	Gly	Asn	Ser	Gly	Cys	Gly		
			420					425					430				
Lys	Ser	Thr	Thr	Val	Gln	Leu	Met	Gln	Arg	Leu	Tyr	Asp	Pro	Thr	Glu		
		435					440					445					
Gly	Met	Val	Ser	Val	Asp	Gly	Gln	Asp	Ile	Arg	Thr	Ile	Asn	Val	Arg		
	450					455					460						
Phe	Leu	Arg	Glu	Ile	Ile	Gly	Val	Val	Ser	Gln	Glu	Pro	Val	Leu	Phe		
465					470					475					480		
Ala	Thr	Thr	Ile	Ala	Glu	Asn	Ile	Arg	Tyr	Gly	Arg	Glu	Asn	Val	Thr		
				485					490					495			
Met	Asp	Glu	Ile	Glu	Lys	Ala	Val	Lys	Glu	Ala	Asn	Ala	Tyr	Asp	Phe		
			500					505					510				
Ile	Met	Lys	Leu	Pro	His	Lys	Phe	Asp	Thr	Leu	Val	Gly	Glu	Arg	Gly		
		515					520					525					
Ala	Gln	Leu	Ser	Gly	Gly	Gln	Lys	Gln	Arg	Ile	Ala	Ile	Ala	Arg	Ala		
	530					535					540						
Leu	Val	Arg	Asn	Pro	Lys	Ile	Leu	Leu	Leu	Asp	Glu	Ala	Thr	Ser	Ala		
545					550					555					560		
Leu	Asp	Thr	Glu	Ser	Glu	Ala	Val	Val	Gln	Val	Ala	Leu	Asp	Lys	Ala		
				565					570					575			
Arg	Lys	Gly	Arg	Thr	Thr	Ile	Val	Ile	Ala	His	Arg	Leu	Ser	Thr	Val		
			580					585					590				
Arg	Asn	Ala	Asp	Val	Ile	Ala	Gly	Phe	Asp	Asp	Gly	Val	Ile	Val	Glu		
		595					600					605					
Lys	Gly	Asn	His	Asp	Glu	Leu	Met	Lys	Glu	Lys	Gly	Ile	Tyr	Phe	Lys		
	610					615					620						
Leu	Val	Thr	Met	Gln	Thr	Ala	Gly	Asn	Glu	Val	Glu	Leu	Glu	Asn	Ala		
625					630					635					640		
Ala	Asp	Glu	Ser	Lys	Ser	Glu	Ile	Asp	Ala	Leu	Glu	Met	Ser	Ser	Asn		
				645					650					655			
Asp	Ser	Arg	Ser	Ser	Leu	Ile	Arg	Lys	Arg	Ser	Thr	Arg	Arg	Ser	Val		
			660					665					670				
Arg	Gly	Ser	Gln	Ala	Gln	Asp	Arg	Lys	Leu	Ser	Thr	Lys	Glu	Ala	Leu		
		675					680					685					
Asp	Glu	Ser	Ile	Pro	Pro	Val	Ser	Phe	Trp	Arg	Ile	Met	Lys	Leu	Asn		
	690					695					700						
Leu	Thr	Glu	Trp	Pro	Tyr	Phe	Val	Val	Gly	Val	Phe	Cys	Ala	Ile	Ile		
705					710					715					720		
Asn	Gly	Gly	Leu	Gln	Pro	Ala	Phe	Ala	Ile	Ile	Phe	Ser	Lys	Ile	Ile		
				725					730					735			
Gly	Val	Phe	Thr	Arg	Ile	Asp	Asp	Pro	Glu	Thr	Lys	Arg	Gln	Asn	Ser		
			740					745					750				

Asn Leu Phe Ser Leu Leu Phe Leu Ala Leu Gly Ile Ile Ser Phe Ile
 755 760 765
 Thr Phe Phe Leu Gln Gly Phe Thr Phe Gly Lys Ala Gly Glu Ile Leu
 770 775 780
 Thr Lys Arg Leu Arg Tyr Met Val Phe Arg Ser Met Leu Arg Gln Asp
 785 790 795 800
 Val Ser Trp Phe Asp Asp Pro Lys Asn Thr Thr Gly Ala Leu Thr Thr
 805 810 815
 Arg Leu Ala Asn Asp Ala Ala Gln Val Lys Gly Ala Ile Gly Ser Arg
 820 825 830
 Leu Ala Val Ile Thr Gln Asn Ile Ala Asn Leu Gly Thr Gly Ile Ile
 835 840 845
 Ile Ser Phe Ile Tyr Gly Trp Gln Leu Thr Leu Leu Leu Ala Ile
 850 855 860
 Val Pro Ile Ile Ala Ile Ala Gly Val Val Glu Met Lys Met Leu Ser
 865 870 875 880
 Gly Gln Ala Leu Lys Asp Lys Lys Glu Leu Glu Gly Ala Gly Lys Ile
 885 890 895
 Ala Thr Glu Ala Ile Glu Asn Phe Arg Thr Val Val Ser Leu Thr Gln
 900 905 910
 Glu Gln Lys Phe Glu His Met Tyr Ala Gln Ser Leu Gln Val Pro Tyr
 915 920 925
 Arg Asn Ser Leu Arg Lys Ala His Ile Phe Gly Ile Thr Phe Ser Phe
 930 935 940
 Thr Gln Ala Met Met Tyr Phe Ser Tyr Ala Gly Cys Phe Arg Phe Gly
 945 950 955 960
 Ala Tyr Leu Val Ala His Lys Leu Met Ser Phe Glu Asp Val Leu Leu
 965 970 975
 Val Phe Ser Ala Val Val Phe Gly Ala Met Ala Val Gly Gln Val Ser
 980 985 990
 Ser Phe Ala Pro Asp Tyr Ala Lys Ala Lys Ile Ser Ala Ala His Ile
 995 1000 1005
 Ile Met Ile Ile Glu Lys Thr Pro Leu Ile Asp Ser Tyr Ser Thr
 1010 1015 1020
 Glu Gly Leu Met Pro Asn Thr Leu Glu Gly Asn Val Thr Phe Gly
 1025 1030 1035
 Glu Val Val Phe Asn Tyr Pro Thr Arg Pro Asp Ile Pro Val Leu
 1040 1045 1050
 Gln Gly Leu Ser Leu Glu Val Lys Lys Gly Gln Thr Leu Ala Leu
 1055 1060 1065
 Val Gly Ser Ser Gly Cys Gly Lys Ser Thr Val Val Gln Leu Leu
 1070 1075 1080
 Glu Arg Phe Tyr Asp Pro Leu Ala Gly Lys Val Leu Leu Asp Gly
 1085 1090 1095

Lys Glu Ile Lys Arg Leu Asn Val Gln Trp Leu Arg Ala His Leu
 1100 1105 1110
 Gly Ile Val Ser Gln Glu Pro Ile Leu Phe Asp Cys Ser Ile Ala
 1115 1120 1125
 Glu Asn Ile Ala Tyr Gly Asp Asn Ser Arg Val Val Ser Gln Glu
 1130 1135 1140
 Glu Ile Val Arg Ala Ala Lys Glu Ala Asn Ile His Ala Phe Ile
 1145 1150 1155
 Glu Ser Leu Pro Asn Lys Tyr Ser Thr Lys Val Gly Asp Lys Gly
 1160 1165 1170
 Thr Gln Leu Ser Gly Gly Gln Lys Gln Arg Ile Ala Ile Ala Arg
 1175 1180 1185
 Ala Leu Val Arg Gln Pro His Ile Leu Leu Leu Asp Glu Ala Thr
 1190 1195 1200
 Ser Ala Leu Asp Thr Glu Ser Glu Lys Val Val Gln Glu Ala Leu
 1205 1210 1215
 Asp Lys Ala Arg Glu Gly Arg Thr Cys Ile Val Ile Ala His Arg
 1220 1225 1230
 Leu Ser Thr Ile Gln Asn Ala Asp Leu Ile Val Val Phe Gln Asn
 1235 1240 1245
 Gly Arg Val Lys Glu His Gly Thr His Gln Gln Leu Leu Ala Gln
 1250 1255 1260
 Lys Gly Ile Tyr Phe Ser Met Val Ser Val Gln Ala Gly Thr Lys
 1265 1270 1275
 Arg Gln
 1280

<210> 5 <211> 4189 <212> DNA <213> Mus musculus <400> 5
 atggagttttg aagagaacct taagggaaga gcagacaaga acttctcgaa gatgggcaaa 60
 aagagtaaaa aggagaagaa agaaaagaaa cctgctgttg gcgtattttg gatgtttcgc 120
 tatgcagatt ggctggacaa gctgtgcatg attctgggaa ctctcgctgc tattatccat 180
 ggaacattac ttcccctctt gatgctggtg tttggaaaca tgacagatag ttttcaaaaa 240
 gcagaagcca gtattctgcc aagcattact aatcaaagtg gacccaacag tactctgac 300
 atcagcaaca gcagtctgga ggaagagatg gccatatacg cctactatta caccgggatt 360
 ggtgctggtg tgctcatagt tgcctacatc caggttttcac tttggtgcct ggcagctgga 420
 agacagatac acaagattag gcagaagttt ttccatgcta taatgaatca ggagataggc 480
 tggtttgatg tgcattgatg tggggagctc aacacccggc tcacagatga tgtctccaaa 540
 attaatgacg gaattggtga caaaattggg atgttttttc agtccataac cacattttta 600
 gccggtttta tcataggatt tataagtggg tggaagctaa cccttgatcat tttggctgtc 660
 agccctctta ttggattgtc atctgctttg tgggcaaagg tattgacttc atttactaat 720
 aaggaactcc aggcttatgc aaaagctgga gcagttgctg aagaagtctt agcagccatc 780

agaactgtga	ttgccttttg	aggacaacag	aaggaacttg	aaaggtacaa	taaaaattta	840
gaagaagcta	aaaatgttgg	cataaagaaa	gctatcacag	ccagcatttc	gataggcatt	900
gcctacctgt	tggctctatgc	atcatatgca	ctggcattct	ggatatgggac	atccttggtc	960
ctctcaaatg	aatattctat	tggagaagtg	cttactgtct	tcttctctat	tttggtgggg	1020
acttttagta	ttggacactt	ggccccaac	atagaagcct	ttgcaaacgc	acgaggggca	1080
gcctttgaaa	tcttcaagat	aattgataac	gagccaagca	ttgacagctt	ctcaacaaag	1140
ggctacaaac	cagacagtat	aatgggaaac	ttagagttta	aaaatgttca	cttcaactac	1200
ccatcgagaa	gcgaagttca	gatcttgaag	ggcctcaatc	tgaaggtgaa	gagcggacag	1260
acggtggcct	tggttggcaa	cagtggctgt	ggaaaaagca	caactgtcca	gctgatgcag	1320
aggctctacg	accccttga	gggcgtggtc	agtatcgacg	gacaagacat	cagaaccatc	1380
aatgtgaggt	atctgagggg	gatcattggg	gtgggtgagtc	aggaacctgt	gctgtttgcc	1440
accacgatcg	ccgagaacat	tcgctatggc	cgagaagatg	tcaccatgga	tgagattgag	1500
aaagctgtca	aggaagccaa	tgcctatgac	ttcatcatga	aactgcccc	ccaatttgac	1560
accctggttg	gtgagagagg	ggcgcagctg	agtgggggac	agaaacagag	aatcgccatt	1620
gccccggccc	tggctccgaa	tccaagatc	cttttgttgg	acgaggccac	ctcagccctg	1680
gatacagaaa	gtgaagctgt	ggtgcaggcc	gcactggata	aggctagaga	aggccggacc	1740
accattgtga	tagctcatcg	cttgtctaca	gttcgtaatg	ctgacgtcat	tgctggtttt	1800
gatggtggtg	tcattgtgga	gcaaggaaat	catgatgagc	tcattgagaga	aaagggcatt	1860
tacttcaaac	ttgtcatgac	acagactaga	ggaaatgaaa	ttgaaccagg	aaataatgct	1920
tatggatccc	agagtgcac	tgatgcttct	gaactgactt	cagaagaatc	caaatcacct	1980
ttaataagga	gatcaattta	cagaagtgtc	cacagaaagc	aagaccaaga	gagaagactt	2040
agtatgaaag	aggctgtgga	tgaagatgtg	cctctggttt	ccttttggcg	gatcctaaat	2100
ctaaatctaa	gtgaatggcc	ttatttactt	gttggcgtag	tttgcgctgt	tataaatggg	2160
tgcatacaac	cagtgtttgc	catagtattt	tcaaggattg	taggggtttt	ttcaagagat	2220
gatgaccatg	aaactaaacg	acagaattgt	aatttgtttt	ccctgttctt	tctggttatg	2280
gggctgattt	cttttgttac	atatttcttt	cagggttca	catttggtgaa	agccggagag	2340
atcctcacca	agcgagtccg	atacatggtt	ttcaaateca	tgctgagaca	ggatataagc	2400
tggttcgatg	accataagaa	cagcactggc	tactgacca	ccaggctcgc	cagtgatgct	2460
tctagtgtta	aagggcgat	gggcgccagg	cttgctgtag	ttaccagaa	tgtagcaaac	2520
ctcgggacag	gagtcattct	ctccttagtc	tatggctggc	agctgacact	tctacttgta	2580
gtaattatac	cgctcattgt	attgggcgga	attattgaaa	tgaagctgtt	gtctggccaa	2640
gccttgaagg	acaagaaaca	gcttgagatc	tctgggaaga	ttgctacaga	agcaattgaa	2700
aacttccgca	ctattgtctc	tttgactcgg	gagcagaagt	ttgaaacat	gtatgccag	2760

agcttgcagg taccatacag aaatgcatg aagaaagcac acgtgttttg gatcacgttc 2820
 tccttcaccc aggccatgat gtatttttct tatgtctgctt gtttcgggtt cgggtgcctac 2880
 ttggtggcac aacaactcat gacttttgaa aatgttatgt tggatatttc tgctgttgctc 2940
 tttggtgcc a tggcagctgg gaatactagt tcatttgctc ctgactatgc gaaagccaaa 3000
 gtatcagcat ctcatatcat caggatcatt gagaaaaccc ctgagattga cagctacagc 3060
 acagaggggt tgaagcctac tctgttagaa ggaaatgtaa aatttaatgg agtccagttt 3120
 aactatccca cccgacccaa catcccagtg cttcaggggc tgagcctcga ggtgaagaag 3180
 ggccagacgt tggccctggt gggcagcagt ggctgtggga agagcacagt ggtccagctg 3240
 ctcgagcgct tctacgaccc catggctgga tcagtgtttc tagatggcaa agaaataaag 3300
 caactgaatg tccagtggct ccgagctcac cttggcattg tgtcccagga gccattctc 3360
 tttgactgca gcattgcaga gaacatcgcc tatggagaca acagccgggc cgtgtctcat 3420
 gaggagattg tgagggcagc caaggaggcc aacatccacc agttcatcga ctactgcct 3480
 gataaatata acaccagagt aggagacaaa ggcactcagc tgtcgggtgg gcagaagcag 3540
 cgcacgcca tcgcacgtgc cctcgtcaga cagcctcaca ttttacttct ggacgaagca 3600
 acatcagctc tggatacaga aagtgaaaag gttgtccagg aagcgctgga caaagccagg 3660
 gaaggccgca cctgcattgt gatcgctcac cgctgtcca ccatccagaa cgcggaactg 3720
 atcgtggtga ttgagaacgg caaagtcaag gagcacggca cccaccagca gctgctggcg 3780
 cagaagggca tctacttctc aatgggccag gctggagcaa agcgctcatg agctgtgact 3840
 atctgaggtg ctaagtattt ttaatatggg tgtttaaaca tggcaccaaa ccaaagttaa 3900
 aaggcaaggg ctgttaaagg taactccatc aagatgagaa gccttcggag actttgtaat 3960
 taaatgaacc aaaatcggaa acaacaaac aaacaaacaa acaagccata gttaaacagg 4020
 gccatgtttt taattgcatt acgtgattca taagagaaca tatagttttt taaaataaaa 4080
 tgtataattt tgtttcagtt ttttaatttct accctacttt cttaaagat tataaagatt 4140
 gtaaaaagca ctatttctta aattgcctat aaaaattaaa ttttcatat 4189

<210> 6 <211> 1276 <212> PRT <213> Mus musculus <400> 6

Met Glu Phe Glu Glu Asn Leu Lys Gly Arg Ala Asp Lys Asn Phe Ser
 1 5 10 15
 Lys Met Gly Lys Lys Ser Lys Lys Glu Lys Lys Glu Lys Lys Pro Ala
 20 25 30
 Val Gly Val Phe Gly Met Phe Arg Tyr Ala Asp Trp Leu Asp Lys Leu
 35 40 45
 Cys Met Ile Leu Gly Thr Leu Ala Ala Ile Ile His Gly Thr Leu Leu
 50 55 60
 Pro Leu Leu Met Leu Val Phe Gly Asn Met Thr Asp Ser Phe Thr Lys
 65 70 75 80

Ala	Glu	Ala	Ser	Ile	Leu	Pro	Ser	Ile	Thr	Asn	Gln	Ser	Gly	Pro	Asn	
				85					90					95		
Ser	Thr	Leu	Ile	Ile	Ser	Asn	Ser	Ser	Leu	Glu	Glu	Glu	Met	Ala	Ile	
			100					105					110			
Tyr	Ala	Tyr	Tyr	Tyr	Thr	Gly	Ile	Gly	Ala	Gly	Val	Leu	Ile	Val	Ala	
		115					120					125				
Tyr	Ile	Gln	Val	Ser	Leu	Trp	Cys	Leu	Ala	Ala	Gly	Arg	Gln	Ile	His	
	130					135					140					
Lys	Ile	Arg	Gln	Lys	Phe	Phe	His	Ala	Ile	Met	Asn	Gln	Glu	Ile	Gly	
145					150					155					160	
Trp	Phe	Asp	Val	His	Asp	Val	Gly	Glu	Leu	Asn	Thr	Arg	Leu	Thr	Asp	
				165					170					175		
Asp	Val	Ser	Lys	Ile	Asn	Asp	Gly	Ile	Gly	Asp	Lys	Ile	Gly	Met	Phe	
			180					185					190			
Phe	Gln	Ser	Ile	Thr	Thr	Phe	Leu	Ala	Gly	Phe	Ile	Ile	Gly	Phe	Ile	
		195					200					205				
Ser	Gly	Trp	Lys	Leu	Thr	Leu	Val	Ile	Leu	Ala	Val	Ser	Pro	Leu	Ile	
	210					215					220					
Gly	Leu	Ser	Ser	Ala	Leu	Trp	Ala	Lys	Val	Leu	Thr	Ser	Phe	Thr	Asn	
225					230					235					240	
Lys	Glu	Leu	Gln	Ala	Tyr	Ala	Lys	Ala	Gly	Ala	Val	Ala	Glu	Glu	Val	
				245					250					255		
Leu	Ala	Ala	Ile	Arg	Thr	Val	Ile	Ala	Phe	Gly	Gly	Gln	Gln	Lys	Glu	
			260					265					270			
Leu	Glu	Arg	Tyr	Asn	Lys	Asn	Leu	Glu	Glu	Ala	Lys	Asn	Val	Gly	Ile	
		275					280					285				
Lys	Lys	Ala	Ile	Thr	Ala	Ser	Ile	Ser	Ile	Gly	Ile	Ala	Tyr	Leu	Leu	
	290					295					300					
Val	Tyr	Ala	Ser	Tyr	Ala	Leu	Ala	Phe	Trp	Tyr	Gly	Thr	Ser	Leu	Val	
305					310					315					320	
Leu	Ser	Asn	Glu	Tyr	Ser	Ile	Gly	Glu	Val	Leu	Thr	Val	Phe	Phe	Ser	
				325					330					335		
Ile	Leu	Leu	Gly	Thr	Phe	Ser	Ile	Gly	His	Leu	Ala	Pro	Asn	Ile	Glu	
			340					345					350			
Ala	Phe	Ala	Asn	Ala	Arg	Gly	Ala	Ala	Phe	Glu	Ile	Phe	Lys	Ile	Ile	
		355					360					365				
Asp	Asn	Glu	Pro	Ser	Ile	Asp	Ser	Phe	Ser	Thr	Lys	Gly	Tyr	Lys	Pro	
	370					375					380					
Asp	Ser	Ile	Met	Gly	Asn	Leu	Glu	Phe	Lys	Asn	Val	His	Phe	Asn	Tyr	
385					390					395					400	
Pro	Ser	Arg	Ser	Glu	Val	Gln	Ile	Leu	Lys	Gly	Leu	Asn	Leu	Lys	Val	
				405					410					415		
Lys	Ser	Gly	Gln	Thr	Val	Ala	Leu	Val	Gly	Asn	Ser	Gly	Cys	Gly	Lys	
			420					425					430			

Ser	Thr	Thr	Val	Gln	Leu	Met	Gln	Arg	Leu	Tyr	Asp	Pro	Leu	Glu	Gly	435	440	445
Val	Val	Ser	Ile	Asp	Gly	Gln	Asp	Ile	Arg	Thr	Ile	Asn	Val	Arg	Tyr	450	455	460
Leu	Arg	Glu	Ile	Ile	Gly	Val	Val	Ser	Gln	Glu	Pro	Val	Leu	Phe	Ala	465	470	475
Thr	Thr	Ile	Ala	Glu	Asn	Ile	Arg	Tyr	Gly	Arg	Glu	Asp	Val	Thr	Met	485	490	495
Asp	Glu	Ile	Glu	Lys	Ala	Val	Lys	Glu	Ala	Asn	Ala	Tyr	Asp	Phe	Ile	500	505	510
Met	Lys	Leu	Pro	His	Gln	Phe	Asp	Thr	Leu	Val	Gly	Glu	Arg	Gly	Ala	515	520	525
Gln	Leu	Ser	Gly	Gly	Gln	Lys	Gln	Arg	Ile	Ala	Ile	Ala	Arg	Ala	Leu	530	535	540
Val	Arg	Asn	Pro	Lys	Ile	Leu	Leu	Leu	Asp	Glu	Ala	Thr	Ser	Ala	Leu	545	550	555
Asp	Thr	Glu	Ser	Glu	Ala	Val	Val	Gln	Ala	Ala	Leu	Asp	Lys	Ala	Arg	565	570	575
Glu	Gly	Arg	Thr	Thr	Ile	Val	Ile	Ala	His	Arg	Leu	Ser	Thr	Val	Arg	580	585	590
Asn	Ala	Asp	Val	Ile	Ala	Gly	Phe	Asp	Gly	Gly	Val	Ile	Val	Glu	Gln	595	600	605
Gly	Asn	His	Asp	Glu	Leu	Met	Arg	Glu	Lys	Gly	Ile	Tyr	Phe	Lys	Leu	610	615	620
Val	Met	Thr	Gln	Thr	Arg	Gly	Asn	Glu	Ile	Glu	Pro	Gly	Asn	Asn	Ala	625	630	635
Tyr	Gly	Ser	Gln	Ser	Asp	Thr	Asp	Ala	Ser	Glu	Leu	Thr	Ser	Glu	Glu	645	650	655
Ser	Lys	Ser	Pro	Leu	Ile	Arg	Arg	Ser	Ile	Tyr	Arg	Ser	Val	His	Arg	660	665	670
Lys	Gln	Asp	Gln	Glu	Arg	Arg	Leu	Ser	Met	Lys	Glu	Ala	Val	Asp	Glu	675	680	685
Asp	Val	Pro	Leu	Val	Ser	Phe	Trp	Arg	Ile	Leu	Asn	Leu	Asn	Leu	Ser	690	695	700
Glu	Trp	Pro	Tyr	Leu	Leu	Val	Gly	Val	Leu	Cys	Ala	Val	Ile	Asn	Gly	705	710	715
Cys	Ile	Gln	Pro	Val	Phe	Ala	Ile	Val	Phe	Ser	Arg	Ile	Val	Gly	Val	725	730	735
Phe	Ser	Arg	Asp	Asp	His	Glu	Thr	Lys	Arg	Gln	Asn	Cys	Asn	Leu		740	745	750
Phe	Ser	Leu	Phe	Phe	Leu	Val	Met	Gly	Leu	Ile	Ser	Phe	Val	Thr	Tyr	755	760	765
Phe	Phe	Gln	Gly	Phe	Thr	Phe	Gly	Lys	Ala	Gly	Glu	Ile	Leu	Thr	Lys	770	775	780

Arg	Val	Arg	Tyr	Met	Val	Phe	Lys	Ser	Met	Leu	Arg	Gln	Asp	Ile	Ser	785	790	795	800
Trp	Phe	Asp	Asp	His	Lys	Asn	Ser	Thr	Gly	Ser	Leu	Thr	Thr	Arg	Leu	805	810	815	
Ala	Ser	Asp	Ala	Ser	Ser	Val	Lys	Gly	Ala	Met	Gly	Ala	Arg	Leu	Ala	820	825	830	
Val	Val	Thr	Gln	Asn	Val	Ala	Asn	Leu	Gly	Thr	Gly	Val	Ile	Leu	Ser	835	840	845	
Leu	Val	Tyr	Gly	Trp	Gln	Leu	Thr	Leu	Leu	Leu	Val	Val	Ile	Ile	Pro	850	855	860	
Leu	Ile	Val	Leu	Gly	Gly	Ile	Ile	Glu	Met	Lys	Leu	Leu	Ser	Gly	Gln	865	870	875	880
Ala	Leu	Lys	Asp	Lys	Lys	Gln	Leu	Glu	Ile	Ser	Gly	Lys	Ile	Ala	Thr	885	890	895	
Glu	Ala	Ile	Glu	Asn	Phe	Arg	Thr	Ile	Val	Ser	Leu	Thr	Arg	Glu	Gln	900	905	910	
Lys	Phe	Glu	Thr	Met	Tyr	Ala	Gln	Ser	Leu	Gln	Val	Pro	Tyr	Arg	Asn	915	920	925	
Ala	Met	Lys	Lys	Ala	His	Val	Phe	Gly	Ile	Thr	Phe	Ser	Phe	Thr	Gln	930	935	940	
Ala	Met	Met	Tyr	Phe	Ser	Tyr	Ala	Ala	Cys	Phe	Arg	Phe	Gly	Ala	Tyr	945	950	955	960
Leu	Val	Ala	Gln	Gln	Leu	Met	Thr	Phe	Glu	Asn	Val	Met	Leu	Val	Phe	965	970	975	
Ser	Ala	Val	Val	Phe	Gly	Ala	Met	Ala	Ala	Gly	Asn	Thr	Ser	Ser	Phe	980	985	990	
Ala	Pro	Asp	Tyr	Ala	Lys	Ala	Lys	Val	Ser	Ala	Ser	His	Ile	Ile	Arg	995	1000	1005	
Ile	Ile	Glu	Lys	Thr	Pro	Glu	Ile	Asp	Ser	Tyr	Ser	Thr	Glu	Gly	1010	1015	1020		
Leu	Lys	Pro	Thr	Leu	Leu	Glu	Gly	Asn	Val	Lys	Phe	Asn	Gly	Val	1025	1030	1035		
Gln	Phe	Asn	Tyr	Pro	Thr	Arg	Pro	Asn	Ile	Pro	Val	Leu	Gln	Gly	1040	1045	1050		
Leu	Ser	Leu	Glu	Val	Lys	Lys	Gly	Gln	Thr	Leu	Ala	Leu	Val	Gly	1055	1060	1065		
Ser	Ser	Gly	Cys	Gly	Lys	Ser	Thr	Val	Val	Gln	Leu	Leu	Glu	Arg	1070	1075	1080		
Phe	Tyr	Asp	Pro	Met	Ala	Gly	Ser	Val	Phe	Leu	Asp	Gly	Lys	Glu	1085	1090	1095		
Ile	Lys	Gln	Leu	Asn	Val	Gln	Trp	Leu	Arg	Ala	His	Leu	Gly	Ile	1100	1105	1110		
Val	Ser	Gln	Glu	Pro	Ile	Leu	Phe	Asp	Cys	Ser	Ile	Ala	Glu	Asn	1115	1120	1125		

Ile	Ala	Tyr	Gly	Asp	Asn	Ser	Arg	Ala	Val	Ser	His	Glu	Glu	Ile
1130						1135					1140			
Val	Arg	Ala	Ala	Lys	Glu	Ala	Asn	Ile	His	Gln	Phe	Ile	Asp	Ser
1145						1150					1155			
Leu	Pro	Asp	Lys	Tyr	Asn	Thr	Arg	Val	Gly	Asp	Lys	Gly	Thr	Gln
1160						1165					1170			
Leu	Ser	Gly	Gly	Gln	Lys	Gln	Arg	Ile	Ala	Ile	Ala	Arg	Ala	Leu
1175						1180					1185			
Val	Arg	Gln	Pro	His	Ile	Leu	Leu	Leu	Asp	Glu	Ala	Thr	Ser	Ala
1190						1195					1200			
Leu	Asp	Thr	Glu	Ser	Glu	Lys	Val	Val	Gln	Glu	Ala	Leu	Asp	Lys
1205						1210					1215			
Ala	Arg	Glu	Gly	Arg	Thr	Cys	Ile	Val	Ile	Ala	His	Arg	Leu	Ser
1220						1225					1230			
Thr	Ile	Gln	Asn	Ala	Asp	Leu	Ile	Val	Val	Ile	Glu	Asn	Gly	Lys
1235						1240					1245			
Val	Lys	Glu	His	Gly	Thr	His	Gln	Gln	Leu	Leu	Ala	Gln	Lys	Gly
1250						1255					1260			
Ile	Tyr	Phe	Ser	Met	Val	Gln	Ala	Gly	Ala	Lys	Arg	Ser		
1265						1270					1275			

<210> 7 <211> 4788 <212> DNA <213> Mus musculus <400> 7

atggaacttg aagaggacct taagggaaga gcagacaaga acttctcaaa gatgggcaaa	60
aagagtaaaa aggagaagaa agaaaagaaa ccagcagtcg gtgtgcttac aatgtttcgt	120
tatgcagggt ggctagacag gttgtacatg ctggtgggaa ctctggctgc tattatccat	180
ggagtggcgc tcccacttat gatgctgac tttggtgaca tgacagatag ctttgcaagt	240
gtaggaaacg tctctaaaaa cagtactaat atgagtgagg ccgataaaag agccatgttt	300
gccaaactgg aggaagaaat gaccacgtac gcctactatt acaccgggat tgggtgctggt	360
gtgctcatag ttgcctacat ccagggtttca ttttgggtgcc tggcagctgg aagacagata	420
cacaagatca ggcagaagtt ttttcatgct ataatgaatc aggagatagg ctggtttgat	480
gtgcatgacg ttggggagct caacaccggg ctacacagatg atgtttccaa aattaatgaa	540
ggaattggtg acaaaatcgg aatgtttctc caggcaatgg caacattttt tgggtggttt	600
ataataggat ttaccctggt ctggaagcta acccttgtga ttttggccat cagccctggt	660
cttggtactgt cagctggtat ttgggcaaag atattgtctt catttactga taaggaactc	720
catgcttatg caaaagctgg agcagttgct gaagaagtct tagcagccat cagaactgtg	780
attgcgtttg gaggacaaaa gaaggaactt gaaaggtaca ataacaactt ggaagaagct	840
aaaaggctgg ggataaagaa agctatcacg gccaacatct ccatgggtgc agctttttctc	900
cttatctatg catcatatgc tctggcattc tgggtatggga cttccttggt catctccaaa	960
gaatactcta ttggacaagt gctcactgct ttcttttccg tgtaattgg agcattcagt	1020

g ttggacagg catctccaaa tattgaagcc ttcgccaatg cacgaggagc agcttatgaa	1080
gtctttcaaaa taattgataa taagcccagt atagacagct tctcaaagag tgggcacaaa	1140
ccagacaaca tacaaggaaa tctggaattt aagaatattc acttcagtta cccatctcga	1200
aaagaagttc agatcttgaa gggcctcaat ctgaagggtga agagcggaca gacggtggcc	1260
ctggttggca acagtggctg tggaaaaagc acaactgtcc agctgatgca aaggctctac	1320
gaccccctag atggcatggt cagtatcgac ggacaggaca tcagaaccat caatgtgagg	1380
tatctgaggg agatcattgg tgtggtgagt caggaacctg tgctgtttgc caccacgac	1440
gccgagaaca ttcgctatgg ccgagaagat gtcaccatgg atgagattga gaaagctgtc	1500
aaggaagcca atgcctatga cttcatcatg aaactgcccc accaatttga caccctgggt	1560
ggtgagagag gggcgcacgt gagtggggga cagaaacaga gaatcgccat tgcccggg	1620
ctggtccgca atcccaagat ctttttggtg gacgaggcca cctcagccct ggatacagaa	1680
agtgaagctg tggttcaggc cgcactggat aaggctagag aaggccggac caccattgtg	1740
atagctcacc gcttgtctac cgttcgtaat gctgacgtca ttgctgggtt tgatgggtgt	1800
gtcattgtgg agcaaggaaa tcatgatgag ctcatgagag aaaagggcat ttacttcaaa	1860
cttgtcatga cacagacagc aggaaatgaa attgaattag gaaatgaagc ttgtaaatct	1920
aaggatgaaa ttgataattt agacatgtct tcaaaagatt caggatccag tctaataaga	1980
agaagatcaa ctgcgaaaag catctgtgga ccacatgacc aagacaggaa gcttagtacc	2040
aaagaggccc tggatgaaga tgtacctcca gcttcctttt ggcggatcct gaagttgaat	2100
tcaactgaat ggccttattt tgtggttggg atattctgtg ccataataaa tggaggctta	2160
cagccagcat tctccgtaat attttcaaaa gttgtagggg tttttacaaa tgggtggccc	2220
cctgaaaccc agcggcagaa cagcaacttg ttttccttgt tgtttctgat ccttgggatc	2280
atttctttca ttacattttt tcttcagggc ttcacatttg gcaaagctgg agagatcctc	2340
accaagcgac tccgatacat ggttttcaaa tccatgctga gacaggatgt gagctgggtt	2400
gatgacccta aaaacaccac cggagcactg accaccaggc tcgccaacga tgctgctcaa	2460
gtgaaagggg ctacagggtc taggcttgct gtgattttcc agaacatagc aaatcttggg	2520
acaggaatca tcatatccct aatctatggc tggcaactaa cacttttact cttagcaatt	2580
gtacccatca ttgcgatagc tggagtgggt gaaatgaaaa tgttgtctgg acaagcactg	2640
aaagataaga aggaactaga aggttctgga aagattgcta cggaagcaat tgaaaacttc	2700
cgcactgttg tctctttgac tcgggagcag aagtttgaaa ccatgtatgc ccagagcttg	2760
cagataccat acagaaatgc gatgaagaaa gcacacgtgt ttgggatcac gttctccttc	2820
accaggcca tgatgtattt ttcttatgct gcttggttcc ggttcggtgc ctacttgggtg	2880
acacaacaac tcatgacttt tgaaaatggt ctggttagtat tctcagctat tgtctttggt	2940
gccatggcag tggggcaggc cagttcattc gctcctgact atgcgaaagc aacagtgtca	3000

```

gcacccaca tcacaggat cattgagaaa acccccgaga ttgacagcta cagcacgcaa 3060
ggcctaaagc cgaatatgtt ggaaggaaat gtgcaattta gtggagtcgt gttcaactat 3120
cccacccgac ccagcatccc agtgcttcag gggctgagcc ttgagggtgaa gaagggccag 3180
acgctggccc tgggtgggcag cagtggctgc gggaagagca cagtgggtcca gctgctcgag 3240
cgcttctacg accccatggc tggatcagtg tttctagatg gcaaagaaat aaagcaactg 3300
aatgtccagt ggctccgagc acagctgggc attgtgtccc aagagcccat tctctttgac 3360
tgcagcatcg cagagaacat tgcctacgga gacaacagcc gggctcgtgtc ttatgaggag 3420
attgtgaggg cagccaagga ggccaacatc caccagttca tcgactcgct acctgataaa 3480
tacaacacca gagtaggaga caaaggcact cagctgtcgg gtgggcagaa gcagcgcac 3540
gccatcgcac gcgccctcgt cagacagcct cacattttac ttctggacga agcaacatca 3600
gctctggata cagaaagtga aaagggtgtc caggaagcgc tggacaaagc caggggaaggc 3660
cgcacctgca ttgtgatcgc tcaccgctg tccaccatcc agaacgcgga cttgatcgtg 3720
gtgattcaga acggcaaggt caaggagcac ggcaccacc agcagctgct ggcgcagaag 3780
ggcatctact tctcaatggg cagtgtgcag gctggagcaa agcgtcatg aactgtgacc 3840
atgtaagatg ttaagtattt ttattgtttg tattcatata tgggtgttta tccaagtcaa 3900
aaggaaaaca cttactaaaa tagccagtta tctattttct gccacagtgg aaagcattta 3960
gtttgggtta gagtcttcag aggctttgta attaaaaaaa caaaaataga tacagcatca 4020
aatggagatt aatgctttta aatgcactat aaaatttata aaagggttaa aagtgaatgt 4080
ttgataatat atacttttat ttatactttc tcatttgtaa ctataactga tttctgctta 4140
acaaattatg tatgtatcaa aaattactga aatgtttgta taaagtatat atagtgaac 4200
tgagcattca tatttttgag ttattttgct caaatgcatg cgaaattata tattgtccca 4260
actgggatat tgtacataat tttagccttt aaaaaacagt ccattactgg ggggaggggg 4320
catcactcta tgggcaaagt gttactcaga catgggcacc tgagttcaga tccctaccac 4380
ctaagtaagc agacaagggt tgggtgtttt gtaatgccag tgctagaggc agaaaagaca 4440
gatcctgcag gctcagtggc tggccaaaca gcctagccaa catagcgcgt tccaggttca 4500
gtgagaaaac ttgtctcaaa aatcagaggg aaaagcaaat gaggtgtcag ccatgtgcac 4560
tcatgcaa at gccatacatg cagaagtatg tgcacacaca cgcacacatt aaccaacgac 4620
tagcaaggaa aatgaagggt gataagaggg gtgggactgg gacaaaggag ggtacctgga 4680
tgaatatgac tgaaggacgt tatgtacaca tatgaaaacg tcgtactgaa actcactaca 4740
atgtatactt aatatattgc taataaaata tttttaaaag aaaaaaat 4788

```

<210> 8 <211> 1276 <212> PRT <213> Mus musculus <400> 8

Met Glu Leu Glu Glu Asp Leu Lys Gly Arg Ala Asp Lys Asn Phe Ser
1 5 10 15

Lys	Met	Gly	Lys	Lys	Ser	Lys	Lys	Glu	Lys	Lys	Glu	Lys	Lys	Pro	Ala		
			20					25						30			
Val	Ser	Val	Leu	Thr	Met	Phe	Arg	Tyr	Ala	Gly	Trp	Leu	Asp	Arg	Leu		
		35					40						45				
Tyr	Met	Leu	Val	Gly	Thr	Leu	Ala	Ala	Ile	Ile	His	Gly	Val	Ala	Leu		
	50					55					60						
Pro	Leu	Met	Met	Leu	Ile	Phe	Gly	Asp	Met	Thr	Asp	Ser	Phe	Ala	Ser		
65					70					75					80		
Val	Gly	Asn	Val	Ser	Lys	Asn	Ser	Thr	Asn	Met	Ser	Glu	Ala	Asp	Lys		
				85					90					95			
Arg	Ala	Met	Phe	Ala	Lys	Leu	Glu	Glu	Glu	Met	Thr	Thr	Tyr	Ala	Tyr		
			100					105						110			
Tyr	Tyr	Thr	Gly	Ile	Gly	Ala	Gly	Val	Leu	Ile	Val	Ala	Tyr	Ile	Gln		
		115					120						125				
Val	Ser	Phe	Trp	Cys	Leu	Ala	Ala	Gly	Arg	Gln	Ile	His	Lys	Ile	Arg		
	130					135					140						
Gln	Lys	Phe	Phe	His	Ala	Ile	Met	Asn	Gln	Glu	Ile	Gly	Trp	Phe	Asp		
145					150					155					160		
Val	His	Asp	Val	Gly	Glu	Leu	Asn	Thr	Arg	Leu	Thr	Asp	Asp	Val	Ser		
				165					170					175			
Lys	Ile	Asn	Glu	Gly	Ile	Gly	Asp	Lys	Ile	Gly	Met	Phe	Phe	Gln	Ala		
			180					185						190			
Met	Ala	Thr	Phe	Phe	Gly	Gly	Phe	Ile	Ile	Gly	Phe	Thr	Arg	Gly	Trp		
		195					200						205				
Lys	Leu	Thr	Leu	Val	Ile	Leu	Ala	Ile	Ser	Pro	Val	Leu	Gly	Leu	Ser		
	210					215					220						
Ala	Gly	Ile	Trp	Ala	Lys	Ile	Leu	Ser	Ser	Phe	Thr	Asp	Lys	Glu	Leu		
225					230					235					240		
His	Ala	Tyr	Ala	Lys	Ala	Gly	Ala	Val	Ala	Glu	Glu	Val	Leu	Ala	Ala		
				245					250					255			
Ile	Arg	Thr	Val	Ile	Ala	Phe	Gly	Gly	Gln	Lys	Lys	Glu	Leu	Glu	Arg		
			260					265						270			
Tyr	Asn	Asn	Asn	Leu	Glu	Glu	Ala	Lys	Arg	Leu	Gly	Ile	Lys	Lys	Ala		
		275					280					285					
Ile	Thr	Ala	Asn	Ile	Ser	Met	Gly	Ala	Ala	Phe	Leu	Leu	Ile	Tyr	Ala		
	290					295					300						
Ser	Tyr	Ala	Leu	Ala	Phe	Trp	Tyr	Gly	Thr	Ser	Leu	Val	Ile	Ser	Lys		
305					310					315					320		
Glu	Tyr	Ser	Ile	Gly	Gln	Val	Leu	Thr	Val	Phe	Phe	Ser	Val	Leu	Ile		
				325					330					335			
Gly	Ala	Phe	Ser	Val	Gly	Gln	Ala	Ser	Pro	Asn	Ile	Glu	Ala	Phe	Ala		
			340					345					350				
Asn	Ala	Arg	Gly	Ala	Ala	Tyr	Glu	Val	Phe	Lys	Ile	Ile	Asp	Asn	Lys		
		355					360					365					

Pro	Ser	Ile	Asp	Ser	Phe	Ser	Lys	Ser	Gly	His	Lys	Pro	Asp	Asn	Ile	370	375	380
Gln	Gly	Asn	Leu	Glu	Phe	Lys	Asn	Ile	His	Phe	Ser	Tyr	Pro	Ser	Arg	385	390	395
Lys	Glu	Val	Gln	Ile	Leu	Lys	Gly	Leu	Asn	Leu	Lys	Val	Lys	Ser	Gly	405	410	415
Gln	Thr	Val	Ala	Leu	Val	Gly	Asn	Ser	Gly	Cys	Gly	Lys	Ser	Thr	Thr	420	425	430
Val	Gln	Leu	Met	Gln	Arg	Leu	Tyr	Asp	Pro	Leu	Asp	Gly	Met	Val	Ser	435	440	445
Ile	Asp	Gly	Gln	Asp	Ile	Arg	Thr	Ile	Asn	Val	Arg	Tyr	Leu	Arg	Glu	450	455	460
Ile	Ile	Gly	Val	Val	Ser	Gln	Glu	Pro	Val	Leu	Phe	Ala	Thr	Thr	Ile	465	470	475
Ala	Glu	Asn	Ile	Arg	Tyr	Gly	Arg	Glu	Asp	Val	Thr	Met	Asp	Glu	Ile	485	490	495
Glu	Lys	Ala	Val	Lys	Glu	Ala	Asn	Ala	Tyr	Asp	Phe	Ile	Met	Lys	Leu	500	505	510
Pro	His	Gln	Phe	Asp	Thr	Leu	Val	Gly	Glu	Arg	Gly	Ala	His	Val	Ser	515	520	525
Gly	Gly	Gln	Lys	Gln	Arg	Ile	Ala	Ile	Ala	Arg	Ala	Leu	Val	Arg	Asn	530	535	540
Pro	Lys	Ile	Leu	Leu	Leu	Asp	Glu	Ala	Thr	Ser	Ala	Leu	Asp	Thr	Glu	545	550	555
Ser	Glu	Ala	Val	Val	Gln	Ala	Ala	Leu	Asp	Lys	Ala	Arg	Glu	Gly	Arg	565	570	575
Thr	Thr	Ile	Val	Ile	Ala	His	Arg	Leu	Ser	Thr	Val	Arg	Asn	Ala	Asp	580	585	590
Val	Ile	Ala	Gly	Phe	Asp	Gly	Gly	Val	Ile	Val	Glu	Gln	Gly	Asn	His	595	600	605
Asp	Glu	Leu	Met	Arg	Glu	Lys	Gly	Ile	Tyr	Phe	Lys	Leu	Val	Met	Thr	610	615	620
Gln	Thr	Ala	Gly	Asn	Glu	Ile	Glu	Leu	Gly	Asn	Glu	Ala	Cys	Lys	Ser	625	630	635
Lys	Asp	Glu	Ile	Asp	Asn	Leu	Asp	Met	Ser	Ser	Lys	Asp	Ser	Gly	Ser	645	650	655
Ser	Leu	Ile	Arg	Arg	Arg	Ser	Thr	Arg	Lys	Ser	Ile	Cys	Gly	Pro	His	660	665	670
Asp	Gln	Asp	Arg	Lys	Leu	Ser	Thr	Lys	Glu	Ala	Leu	Asp	Glu	Asp	Val	675	680	685
Pro	Pro	Ala	Ser	Phe	Trp	Arg	Ile	Leu	Lys	Leu	Asn	Ser	Thr	Glu	Trp	690	695	700
Pro	Tyr	Phe	Val	Val	Gly	Ile	Phe	Cys	Ala	Ile	Ile	Asn	Gly	Gly	Leu	705	710	715

Gln	Pro	Ala	Phe	Ser	Val	Ile	Phe	Ser	Lys	Val	Val	Gly	Val	Phe	Thr		
				725					730					735			
Asn	Gly	Gly	Pro	Pro	Glu	Thr	Gln	Arg	Gln	Asn	Ser	Asn	Leu	Phe	Ser		
			740					745					750				
Leu	Leu	Phe	Leu	Ile	Leu	Gly	Ile	Ile	Ser	Phe	Ile	Thr	Phe	Phe	Leu		
		755					760					765					
Gln	Gly	Phe	Thr	Phe	Gly	Lys	Ala	Gly	Glu	Ile	Leu	Thr	Lys	Arg	Leu		
	770					775					780						
Arg	Tyr	Met	Val	Phe	Lys	Ser	Met	Leu	Arg	Gln	Asp	Val	Ser	Trp	Phe		
785					790					795					800		
Asp	Asp	Pro	Lys	Asn	Thr	Thr	Gly	Ala	Leu	Thr	Thr	Arg	Leu	Ala	Asn		
				805					810					815			
Asp	Ala	Ala	Gln	Val	Lys	Gly	Ala	Thr	Gly	Ser	Arg	Leu	Ala	Val	Ile		
			820					825					830				
Phe	Gln	Asn	Ile	Ala	Asn	Leu	Gly	Thr	Gly	Ile	Ile	Ile	Ser	Leu	Ile		
		835					840					845					
Tyr	Gly	Trp	Gln	Leu	Thr	Leu	Leu	Leu	Leu	Ala	Ile	Val	Pro	Ile	Ile		
	850					855					860						
Ala	Ile	Ala	Gly	Val	Val	Glu	Met	Lys	Met	Leu	Ser	Gly	Gln	Ala	Leu		
865					870					875					880		
Lys	Asp	Lys	Lys	Glu	Leu	Glu	Gly	Ser	Gly	Lys	Ile	Ala	Thr	Glu	Ala		
				885					890					895			
Ile	Glu	Asn	Phe	Arg	Thr	Val	Val	Ser	Leu	Thr	Arg	Glu	Gln	Lys	Phe		
		900						905					910				
Glu	Thr	Met	Tyr	Ala	Gln	Ser	Leu	Gln	Ile	Pro	Tyr	Arg	Asn	Ala	Met		
		915					920					925					
Lys	Lys	Ala	His	Val	Phe	Gly	Ile	Thr	Phe	Ser	Phe	Thr	Gln	Ala	Met		
		930				935					940						
Met	Tyr	Phe	Ser	Tyr	Ala	Ala	Cys	Phe	Arg	Phe	Gly	Ala	Tyr	Leu	Val		
945					950					955					960		
Thr	Gln	Gln	Leu	Met	Thr	Phe	Glu	Asn	Val	Leu	Leu	Val	Phe	Ser	Ala		
				965				970						975			
Ile	Val	Phe	Gly	Ala	Met	Ala	Val	Gly	Gln	Val	Ser	Ser	Phe	Ala	Pro		
			980					985					990				
Asp	Tyr	Ala	Lys	Ala	Thr	Val	Ser	Ala	Ser	His	Ile	Ile	Arg	Ile	Ile		
		995					1000					1005					
Glu	Lys	Thr	Pro	Glu	Ile	Asp	Ser	Tyr	Ser	Thr	Gln	Gly	Leu	Lys			
	1010					1015					1020						
Pro	Asn	Met	Leu	Glu	Gly	Asn	Val	Gln	Phe	Ser	Gly	Val	Val	Phe			
	1025					1030					1035						
Asn	Tyr	Pro	Thr	Arg	Pro	Ser	Ile	Pro	Val	Leu	Gln	Gly	Leu	Ser			
	1040					1045					1050						
Leu	Glu	Val	Lys	Lys	Gly	Gln	Thr	Leu	Ala	Leu	Val	Gly	Ser	Ser			
	1055					1060					1065						

Gly	Cys	Gly	Lys	Ser	Thr	Val	Val	Gln	Leu	Leu	Glu	Arg	Phe	Tyr
1070						1075					1080			
Asp	Pro	Met	Ala	Gly	Ser	Val	Phe	Leu	Asp	Gly	Lys	Glu	Ile	Lys
1085						1090					1095			
Gln	Leu	Asn	Val	Gln	Trp	Leu	Arg	Ala	Gln	Leu	Gly	Ile	Val	Ser
1100						1105					1110			
Gln	Glu	Pro	Ile	Leu	Phe	Asp	Cys	Ser	Ile	Ala	Glu	Asn	Ile	Ala
1115						1120					1125			
Tyr	Gly	Asp	Asn	Ser	Arg	Val	Val	Ser	Tyr	Glu	Glu	Ile	Val	Arg
1130						1135					1140			
Ala	Ala	Lys	Glu	Ala	Asn	Ile	His	Gln	Phe	Ile	Asp	Ser	Leu	Pro
1145						1150					1155			
Asp	Lys	Tyr	Asn	Thr	Arg	Val	Gly	Asp	Lys	Gly	Thr	Gln	Leu	Ser
1160						1165					1170			
Gly	Gly	Gln	Lys	Gln	Arg	Ile	Ala	Ile	Ala	Arg	Ala	Leu	Val	Arg
1175						1180					1185			
Gln	Pro	His	Ile	Leu	Leu	Leu	Asp	Glu	Ala	Thr	Ser	Ala	Leu	Asp
1190						1195					1200			
Thr	Glu	Ser	Glu	Lys	Val	Val	Gln	Glu	Ala	Leu	Asp	Lys	Ala	Arg
1205						1210					1215			
Glu	Gly	Arg	Thr	Cys	Ile	Val	Ile	Ala	His	Arg	Leu	Ser	Thr	Ile
1220						1225					1230			
Gln	Asn	Ala	Asp	Leu	Ile	Val	Val	Ile	Gln	Asn	Gly	Lys	Val	Lys
1235						1240					1245			
Glu	His	Gly	Thr	His	Gln	Gln	Leu	Leu	Ala	Gln	Lys	Gly	Ile	Tyr
1250						1255					1260			
Phe	Ser	Met	Val	Ser	Val	Gln	Ala	Gly	Ala	Lys	Arg	Ser		
1265						1270					1275			

<210> 9 <211> 2719 <212> DNA <213> Homo sapiens <400> 9

tttaggaacg caccgtgcac atgcttggtg gtcttgtaa gtggaaactg ctgctttaga	60
gtttgttttg aagggtccggg tgactcatcc caacatttac atccttaatt gttaaagcgc	120
tgctccgag cgcacgcac ctgagatcct gagcctttgg ttaagaccga gctctattaa	180
gctgaaaaga taaaaactct ccagatgtct tccagtaatg tcgaagtgtt tatcccagtg	240
tcacaaggaa acaccaatgg cttccccgcg acagtttcca atgacctgaa ggcatttact	300
gaaggagctg tgtaagttt tcataacatc tgctatcgag taaaactgaa gagtggcttt	360
ctaccttgtc gaaaaccagt tgagaaagaa atattatcga atatcaatgg gatcatgaaa	420
cctgggtctca acgccatcct gggaccaca ggtggaggca aatcttcgtt attagatgtc	480
ttagctgcaa ggaaagatcc aagtggatta tctggagatg ttctgataaa tggagcaccg	540
cgacctgcca atttcaaagt taattcaggt tacgtggtac aagatgatgt tgtgatgggc	600
actctgacgg tgagagaaaa cttacagttc tcagcagctc ttcggcttgc aacaactatg	660
acgaatcatg aaaaaaacga acggattaac agggtcattg aagagttagg tctggataaa	720

gtggcagact	ccaaggttgg	aactcagttt	atccgtggtg	tgtctggagg	agaaagaaaa	780
aggactagta	taggaatgga	gcttatcact	gacccctcca	tcttgccctt	ggatgagcct	840
acaactggct	tagactcaag	cacagcaaat	gctgtccctt	tgctcctgaa	aaggatgtct	900
aagcagggac	gaacaatcat	cttctccatt	catcagcctc	gatattccat	cttcaagttg	960
tttgatagcc	tcaccttatt	ggcctcagga	agacttatgt	tccacggggc	tgctcaggag	1020
gccttgggat	actttgaatc	agctgggtat	cactgtgagg	cctataataa	ccctgcagac	1080
ttcttcttgg	acatcattaa	tggagattcc	actgctgtgg	cattaaacag	agaagaagac	1140
tttaaagcca	cagagatcat	agagccttcc	aagcaggata	agccactcat	agaaaaatta	1200
gcggagattt	atgtcaactc	ctccttctac	aaagagacaa	aagctgaatt	acatcaactt	1260
tccgggggtg	agaagaagaa	gaagatcaca	gtcttcaagg	agatcagcta	caccacctcc	1320
ttctgtcatc	aactcagatg	ggtttccaag	cgttcattca	aaaacttgct	gggtaatccc	1380
caggcctcta	tagctcagat	cattgtcaca	gtcgtactgg	gactgggtat	aggtgccatt	1440
tactttgggc	taaaaaatga	ttctactgga	atccagaaca	gagctggggg	tctcttcttc	1500
ctgacgacca	accagtgttt	cagcagtgtt	tcagccgtgg	aactctttgt	ggtagagaag	1560
aagctcttca	tacatgaata	catcagcgga	tactacagag	tgatcatctta	tttcttggga	1620
aaactgttat	ctgatttatt	acccatgagg	atgttaccaa	gtattatatt	tacctgtata	1680
gtgtacttca	tgttaggatt	gaagccaaag	gcagatgcct	tcttcgttat	gatgtttacc	1740
cttatgatgg	tggtttattc	agccagttcc	atggcactgg	ccatagcagc	aggtcagagt	1800
gtggtttctg	tagcaacact	tctcatgacc	atctgttttg	tgtttatgat	gattttttca	1860
ggtctgttgg	tcaatctcac	aaccattgca	tcttggctgt	catggcttca	gtacttcagc	1920
attccacgat	atggatttac	ggctttgcag	cataatgaat	ttttgggaca	aaacttctgc	1980
ccaggactca	atgcaacagg	aaacaatcct	tgtaactatg	caacatgtac	tggcgaagaa	2040
tattttggtaa	agcagggcat	cgatctctca	ccctggggct	tgtggaagaa	tcacgtggcc	2100
ttggcttgta	tgattgttat	tttctcaca	attgcctacc	tgaaattgtt	atttcttaaa	2160
aaatattctt	aaatttcccc	ttaattcagt	atgattttatc	ctcacataaa	aaagaagcac	2220
tttgattgaa	gtattcaatc	aagttttttt	gttgttttct	gttcccttgc	catcacactg	2280
ttgcacagca	gcaattgttt	taaagagata	cattttttaga	aatcacaaca	aactgaatta	2340
aacatgaaag	aacccaagac	atcatgtatc	gcataattagt	taatctcctc	agacagtaac	2400
catggggaag	aaatctggtc	taatttatta	atctaaaaaa	ggagaattga	attctggaaa	2460
ctcctgacaa	gttattactg	tctctggcat	ttgtttctct	atcttttaaa	tgaataggta	2520
ggttagtagc	ccttcagtct	taatacttta	tgatgctatg	gtttgccatt	atttaatat	2580
tgacaaatgt	attaatgcta	tactggaaat	gtaaaattga	aaatatgttg	gaaaaaagat	2640
tctgtcttat	agggtaaaaa	aagccaccgg	tgatagaaaa	aaaatctttt	tgataagcac	2700

attaaagtta atagaactt

2719

```

<210> 10 <211> 655 <212> PRT <213> Homo sapien <400> 10
Met Ser Ser Ser Asn Val Glu Val Phe Ile Pro Val Ser Gln Gly Asn
1          5          10          15
Thr Asn Gly Phe Pro Ala Thr Val Ser Asn Asp Leu Lys Ala Phe Thr
          20          25          30
Glu Gly Ala Val Leu Ser Phe His Asn Ile Cys Tyr Arg Val Lys Leu
          35          40          45
Lys Ser Gly Phe Leu Pro Cys Arg Lys Pro Val Glu Lys Glu Ile Leu
          50          55          60
Ser Asn Ile Asn Gly Ile Met Lys Pro Gly Leu Asn Ala Ile Leu Gly
65          70          75          80
Pro Thr Gly Gly Gly Lys Ser Ser Leu Leu Asp Val Leu Ala Ala Arg
          85          90          95
Lys Asp Pro Ser Gly Leu Ser Gly Asp Val Leu Ile Asn Gly Ala Pro
          100          105          110
Arg Pro Ala Asn Phe Lys Cys Asn Ser Gly Tyr Val Val Gln Asp Asp
          115          120          125
Val Val Met Gly Thr Leu Thr Val Arg Glu Asn Leu Gln Phe Ser Ala
          130          135          140
Ala Leu Arg Leu Ala Thr Thr Met Thr Asn His Glu Lys Asn Glu Arg
145          150          155          160
Ile Asn Arg Val Ile Glu Glu Leu Gly Leu Asp Lys Val Ala Asp Ser
          165          170          175
Lys Val Gly Thr Gln Phe Ile Arg Gly Val Ser Gly Gly Glu Arg Lys
          180          185          190
Arg Thr Ser Ile Gly Met Glu Leu Ile Thr Asp Pro Ser Ile Leu Ser
          195          200          205
Leu Asp Glu Pro Thr Thr Gly Leu Asp Ser Ser Thr Ala Asn Ala Val
          210          215          220
Leu Leu Leu Leu Lys Arg Met Ser Lys Gln Gly Arg Thr Ile Ile Phe
225          230          235          240
Ser Ile His Gln Pro Arg Tyr Ser Ile Phe Lys Leu Phe Asp Ser Leu
          245          250          255
Thr Leu Leu Ala Ser Gly Arg Leu Met Phe His Gly Pro Ala Gln Glu
          260          265          270
Ala Leu Gly Tyr Phe Glu Ser Ala Gly Tyr His Cys Glu Ala Tyr Asn
          275          280          285
Asn Pro Ala Asp Phe Phe Leu Asp Ile Ile Asn Gly Asp Ser Thr Ala
          290          295          300
Val Ala Leu Asn Arg Glu Glu Asp Phe Lys Ala Thr Glu Ile Ile Glu
305          310          315          320

```

Pro Ser Lys Gln Asp Lys Pro Leu Ile Glu Lys Leu Ala Glu Ile Tyr
 325 330 335
 Val Asn Ser Ser Phe Tyr Lys Glu Thr Lys Ala Glu Leu His Gln Leu
 340 345 350
 Ser Gly Gly Glu Lys Lys Lys Lys Ile Thr Val Phe Lys Glu Ile Ser
 355 360 365
 Tyr Thr Thr Ser Phe Cys His Gln Leu Arg Trp Val Ser Lys Arg Ser
 370 375 380
 Phe Lys Asn Leu Leu Gly Asn Pro Gln Ala Ser Ile Ala Gln Ile Ile
 385 390 395 400
 Val Thr Val Val Leu Gly Leu Val Ile Gly Ala Ile Tyr Phe Gly Leu
 405 410 415
 Lys Asn Asp Ser Thr Gly Ile Gln Asn Arg Ala Gly Val Leu Phe Phe
 420 425 430
 Leu Thr Thr Asn Gln Cys Phe Ser Ser Val Ser Ala Val Glu Leu Phe
 435 440 445
 Val Val Glu Lys Lys Leu Phe Ile His Glu Tyr Ile Ser Gly Tyr Tyr
 450 455 460
 Arg Val Ser Ser Tyr Phe Leu Gly Lys Leu Leu Ser Asp Leu Leu Pro
 465 470 475 480
 Met Arg Met Leu Pro Ser Ile Ile Phe Thr Cys Ile Val Tyr Phe Met
 485 490 495
 Leu Gly Leu Lys Pro Lys Ala Asp Ala Phe Phe Val Met Met Phe Thr
 500 505 510
 Leu Met Met Val Ala Tyr Ser Ala Ser Ser Met Ala Leu Ala Ile Ala
 515 520 525
 Ala Gly Gln Ser Val Val Ser Val Ala Thr Leu Leu Met Thr Ile Cys
 530 535 540
 Phe Val Phe Met Met Ile Phe Ser Gly Leu Leu Val Asn Leu Thr Thr
 545 550 555 560
 Ile Ala Ser Trp Leu Ser Trp Leu Gln Tyr Phe Ser Ile Pro Arg Tyr
 565 570 575
 Gly Phe Thr Ala Leu Gln His Asn Glu Phe Leu Gly Gln Asn Phe Cys
 580 585 590
 Pro Gly Leu Asn Ala Thr Gly Asn Asn Pro Cys Asn Tyr Ala Thr Cys
 595 600 605
 Thr Gly Glu Glu Tyr Leu Val Lys Gln Gly Ile Asp Leu Ser Pro Trp
 610 615 620
 Gly Leu Trp Lys Asn His Val Ala Leu Ala Cys Met Ile Val Ile Phe
 625 630 635 640
 Leu Thr Ile Ala Tyr Leu Lys Leu Leu Phe Leu Lys Lys Tyr Ser
 645 650 655

<210> 11 <211> 502 <212> DNA <213> Mus musculus <400> 11
 ttccggcctag gggccgaggc ttatacggcc agttccatgg cactggccat agccacaggc

caaagtgtgg tgtctgtagc aacactactc atgacaatcg cttttgtatt tatgatgctc 120
 ttttctggcc tcttggtgaa tctcagaacc attgggcctt ggctgtcctg gcttcagtag 180
 ttttagcattc ctcatatagg cttcacagct ttgcagtata atgaattctt gggacaagag 240
 ttttgtccag gattcaatgt aacggacaac agcacttgtg ttaacagcta tgcaatatgt 300
 actggtaacg agtacttgat aaatcagggc atcgaactgt caccttgggg actgtggaag 360
 aatcatgtgg ccctggcctg tatgattatt atcttctca caattgccta cctgaaattg 420
 ttgtttctta aaaagtattc ttaatttccc ctttaacgga ctattaattg tactccaatt 480
 aaatatgggc acttttgatta cc 502

<210> 12 <211> 147 <212> PRT <213> Mus musculus <400> 12
 Phe Gly Leu Gly Ala Glu Ala Tyr Thr Ala Ser Ser Met Ala Leu Ala
 1 5 10 15
 Ile Ala Thr Gly Gln Ser Val Val Ser Val Ala Thr Leu Leu Met Thr
 20 25 30
 Ile Ala Phe Val Phe Met Met Leu Phe Ser Gly Leu Leu Val Asn Leu
 35 40 45
 Arg Thr Ile Gly Pro Trp Leu Ser Trp Leu Gln Tyr Phe Ser Ile Pro
 50 55 60
 Arg Tyr Gly Phe Thr Ala Leu Gln Tyr Asn Glu Phe Leu Gly Gln Glu
 65 70 75 80
 Phe Cys Pro Gly Phe Asn Val Thr Asp Asn Ser Thr Cys Val Asn Ser
 85 90 95
 Tyr Ala Ile Cys Thr Gly Asn Glu Tyr Leu Ile Asn Gln Gly Ile Glu
 100 105 110
 Leu Ser Pro Trp Gly Leu Trp Lys Asn His Val Ala Leu Ala Cys Met
 115 120 125
 Ile Ile Ile Phe Leu Thr Ile Ala Tyr Leu Lys Leu Leu Phe Leu Lys
 130 135 140
 Lys Tyr Ser
 145

<210> 13 <211> 2025 <212> DNA <213> Mus musculus <400> 13
 aaaggcataa atcctaaaga tgtcttccag taatgaccac gtgttagtac caatgtcgca 60
 gagaaacaac aacggccttc ctaggatgaa ctccagagcc gttaggacgc tcgcagaagg 120
 agatgtgttg agttttcatc acatcaccta tcgagtgaag gtaaagagtg ggtttctagt 180
 ccggaaaaca gttgagaaag aaatactatc agatatcaat gggatcatga aacctggcct 240
 taatgctatt ctgggaccca caggcggagg caagtcttcg ttgctagatg tcttagcagc 300
 aaggaaagat ccaaagggat tatctggaga tgttttgata aatggagcac ctcaacctgc 360
 ccatttcaaa tgctgttcag gttatgtggt tcaagatgac gttgtgatgg gcaccctgac 420
 agtgagagaa aacttacagt tctcagcagc tcttcgactt ccaacaacta tgaagaatca 480

```

tgaaaaaaat gaacggatta acacaatcat taaagagtta ggtctggaaa aagtagcaga      540
ttctaaggtc ggaactcagt ttatccgtgg catctctgga ggagaaagaa aaaggacaag      600
cataggggatg gagctgatca ctgacccttc catcctcttc ctggatgagc ccacgactgg      660
tttgactca agcacagega atgctgtcct tttgctcctg aaaaggatgt ctaaacaggg      720
tcgaacaatc atcttctcca ttcatcagcc tcggtattcc atctttaagt tgtttgacag      780
cctcacetta ctggcttccg ggaaactcgt gttccatggg ccagcacaga aggccttgga      840
gtactttgca tcagcagggt accactgtga gccctacaac aaccctgcgg attttttcct      900
tgatgtcatc aatggagatt cttctgtctg gatgttaaag agagaggaac aagacaatga      960
agcaaacaag actgaagagc cttccaaggg agagaagcca gtaatagaaa atttatctga     1020
gttttatatc aactctgcca tctatggaga aacaaaagct gaattagatc aacttccagg     1080
agctcaggaa aagaaaggaa catcggcctt caaagagcca gtctatgtta cctctttctg     1140
tcaccagctc cgatggattg ccaggcgctc atttaaaaac ttgctcggga accctcaagc     1200
ttctgttgct cagttaattg ttacagtcac actggggcctt attattggtg ccatttactt     1260
tgatctgaaa tatgatgccg ctggaatgca aaatagagct ggagttttgt ttttcctgac     1320
taccaaccag tgtttttcca gtgtgtcagc tgtggagctg ttcgtagtgg agaagaaact     1380
cttcatacat gagtacatca gtggatatta cagagtgtct tcttacttct ttggaaagggt     1440
gatgtctgat ttactcccca tgaggttcct gccaaagtgt atattcactt gtatattata     1500
cttcagtgtta ggactgaaga agacgggtgga tgcttttttc atcatgatgt ttacccttat     1560
aatggtggct tatacggcca gttccatggc actggccata gccacaggcc aaagtgtggt     1620
gtctgtagca acacttctca tgacaatcgc ttttgatatt atgatgctct tttctggcct     1680
cttggtgaat ctcagaacca ttgggccttg gctgtcctgg cttcagtact ttagcattcc     1740
tcgatatggc ttcacagctt tgcaagtata tgaattcttg ggacaagagt tttgtccagg     1800
attcaatgta acggacaaca gcacttgtgt taacagctat gcaatatgta ctggtaacga     1860
gtacttgata aatcagggca tcgaactgtc accttgggga ctgtggaaga atcatgtggc     1920
cctggcttgt atgattatta tcttcctcac aattgcctac ctgaaattgt tgtttcttaa     1980
aaagtattct taatttcccc tttaacggac tattaattgt actcc                        2025

```

```

<210> 14 <211> 657 <212> PRT <213> Mus musculus <400> 14
Met Ser Ser Ser Asn Asp His Val Leu Val Pro Met Ser Gln Arg Asn
1          5          10          15
Asn Asn Gly Leu Pro Arg Met Asn Ser Arg Ala Val Arg Thr Leu Ala
          20          25          30
Glu Gly Asp Val Leu Ser Phe His His Ile Thr Tyr Arg Val Lys Val
          35          40          45
Lys Ser Gly Phe Leu Val Arg Lys Thr Val Glu Lys Glu Ile Leu Ser
          50          55          60

```

Asp	Ile	Asn	Gly	Ile	Met	Lys	Pro	Gly	Leu	Asn	Ala	Ile	Leu	Gly	Pro	65	70	75	80
Thr	Gly	Gly	Gly	Lys	Ser	Ser	Leu	Leu	Asp	Val	Leu	Ala	Ala	Arg	Lys	85	90	95	
Asp	Pro	Lys	Gly	Leu	Ser	Gly	Asp	Val	Leu	Ile	Asn	Gly	Ala	Pro	Gln	100	105	110	
Pro	Ala	His	Phe	Lys	Cys	Cys	Ser	Gly	Tyr	Val	Val	Gln	Asp	Asp	Val	115	120	125	
Val	Met	Gly	Thr	Leu	Thr	Val	Arg	Glu	Asn	Leu	Gln	Phe	Ser	Ala	Ala	130	135	140	
Leu	Arg	Leu	Pro	Thr	Thr	Met	Lys	Asn	His	Glu	Lys	Asn	Glu	Arg	Ile	145	150	155	160
Asn	Thr	Ile	Ile	Lys	Glu	Leu	Gly	Leu	Glu	Lys	Val	Ala	Asp	Ser	Lys	165	170	175	
Val	Gly	Thr	Gln	Phe	Ile	Arg	Gly	Ile	Ser	Gly	Gly	Glu	Arg	Lys	Arg	180	185	190	
Thr	Ser	Ile	Gly	Met	Glu	Leu	Ile	Thr	Asp	Pro	Ser	Ile	Leu	Phe	Leu	195	200	205	
Asp	Glu	Pro	Thr	Thr	Gly	Leu	Asp	Ser	Ser	Thr	Ala	Asn	Ala	Val	Leu	210	215	220	
Leu	Leu	Leu	Lys	Arg	Met	Ser	Lys	Gln	Gly	Arg	Thr	Ile	Ile	Phe	Ser	225	230	235	240
Ile	His	Gln	Pro	Arg	Tyr	Ser	Ile	Phe	Lys	Leu	Phe	Asp	Ser	Leu	Thr	245	250	255	
Leu	Leu	Ala	Ser	Gly	Lys	Leu	Val	Phe	His	Gly	Pro	Ala	Gln	Lys	Ala	260	265	270	
Leu	Glu	Tyr	Phe	Ala	Ser	Ala	Gly	Tyr	His	Cys	Glu	Pro	Tyr	Asn	Asn	275	280	285	
Pro	Ala	Asp	Phe	Phe	Leu	Asp	Val	Ile	Asn	Gly	Asp	Ser	Ser	Ala	Val	290	295	300	
Met	Leu	Asn	Arg	Glu	Glu	Gln	Asp	Asn	Glu	Ala	Asn	Lys	Thr	Glu	Glu	305	310	315	320
Pro	Ser	Lys	Gly	Glu	Lys	Pro	Val	Ile	Glu	Asn	Leu	Ser	Glu	Phe	Tyr	325	330	335	
Ile	Asn	Ser	Ala	Ile	Tyr	Gly	Glu	Thr	Lys	Ala	Glu	Leu	Asp	Gln	Leu	340	345	350	
Pro	Gly	Ala	Gln	Glu	Lys	Lys	Gly	Thr	Ser	Ala	Phe	Lys	Glu	Pro	Val	355	360	365	
Tyr	Val	Thr	Ser	Phe	Cys	His	Gln	Leu	Arg	Trp	Ile	Ala	Arg	Arg	Ser	370	375	380	
Phe	Lys	Asn	Leu	Leu	Gly	Asn	Pro	Gln	Ala	Ser	Val	Ala	Gln	Leu	Ile	385	390	395	400
Val	Thr	Val	Ile	Leu	Gly	Leu	Ile	Ile	Gly	Ala	Ile	Tyr	Phe	Asp	Leu	405	410	415	

Lys Tyr Asp Ala Ala Gly Met Gln Asn Arg Ala Gly Val Leu Phe Phe
 420 425 430
 Leu Thr Thr Asn Gln Cys Phe Ser Ser Val Ser Ala Val Glu Leu Phe
 435 440 445
 Val Val Glu Lys Lys Leu Phe Ile His Glu Tyr Ile Ser Gly Tyr Tyr
 450 455 460
 Arg Val Ser Ser Tyr Phe Phe Gly Lys Val Met Ser Asp Leu Leu Pro
 465 470 475 480
 Met Arg Phe Leu Pro Ser Val Ile Phe Thr Cys Ile Leu Tyr Phe Met
 485 490 495
 Leu Gly Leu Lys Lys Thr Val Asp Ala Phe Phe Ile Met Met Phe Thr
 500 505 510
 Leu Ile Met Val Ala Tyr Thr Ala Ser Ser Met Ala Leu Ala Ile Ala
 515 520 525
 Thr Gly Gln Ser Val Val Ser Val Ala Thr Leu Leu Met Thr Ile Ala
 530 535 540
 Phe Val Phe Met Met Leu Phe Ser Gly Leu Leu Val Asn Leu Arg Thr
 545 550 555 560
 Ile Gly Pro Trp Leu Ser Trp Leu Gln Tyr Phe Ser Ile Pro Arg Tyr
 565 570 575
 Gly Phe Thr Ala Leu Gln Tyr Asn Glu Phe Leu Gly Gln Glu Phe Cys
 580 585 590
 Pro Gly Phe Asn Val Thr Asp Asn Ser Thr Cys Val Asn Ser Tyr Ala
 595 600 605
 Ile Cys Thr Gly Asn Glu Tyr Leu Ile Asn Gln Gly Ile Glu Leu Ser
 610 615 620
 Pro Trp Gly Leu Trp Lys Asn His Val Ala Leu Ala Cys Met Ile Ile
 625 630 635 640
 Ile Phe Leu Thr Ile Ala Tyr Leu Lys Leu Leu Phe Leu Lys Lys Tyr
 645 650 655
 Ser

<210> 15 <211> 20 <212> DNA <213> Artificial <220> <223> primer

<400> 15
 ccacgctcagc cttggacaca 20

<210> 16 <211> 20 <212> DNA <213> Artificial <220> <223> primer

<400> 16
 gccgcttggt gaggatctct 20

<210> 17 <211> 20 <212> DNA <213> Artificial <220> <223> primer

<400> 17
 ccatagccac aggccaaagt 20

<210> 18 <211> 20 <212> DNA <213> Artificial <220> <223> primer
 <400> 18
 gggccacatg attcttccac 20

<210> 19 <211> 20 <212> DNA <213> Artificial <220> <223> primer
 <400> 19
 ggcctcagga agacttatgt 20

<210> 20 <211> 20 <212> DNA <213> Artificial <220> <223> primer
 <400> 20
 aaggaggtgg tgtagctgat 20

<210> 21 <211> 19 <212> DNA <213> Artificial <220> <223> primer
 <400> 21
 agctggagag atcctcacc 19

<210> 22 <211> 19 <212> DNA <213> Artificial <220> <223> primer
 <400> 22
 agccggagag atcctcacc 19

<210> 23 <211> 20 <212> DNA <213> artificial <220> <223> primer
 <400> 23
 ctgtagctgt caatctcggg 20

<210> 24 <211> 20 <212> DNA <213> Artificial <220> <223> primer
 <400> 24
 ctgtagctgt caatctcagg 20

<210> 25 <211> 20 <212> DNA <213> Artificial <220> <223> primer
 <400> 25
 ctgtagctgt caatcagagg 20

<210> 26 <211> 2247 <212> DNA <213> Homo sapiens <400> 26
 accgtgcaca tgcttggtgg tcttgtaag tggaaactgc tgctttagag tttgtttgga 60
 aggtccgggt gactcatccc aacatttaca tccttaattg ttaaagcgct gcctccgagc 120
 gcacgcatcc tgagatcctg agcctttggt taagaccgag ctctattaag ctgaaaagat 180
 aaaaactctc cagatgtctt ccagtaatgt cgaagttttt atcccagtggt cacaaggaaa 240
 caccaatggc ttccccgcga cagcttccaa tgacctgaag gcatttactg aaggagctgt 300
 gttaagtttt cataacatct gctatcgagt aaaactgaag agtggctttc taccttgtag 360
 aaaaccagtt gagaaagaaa tattatcgaa tatcaatggg atcatgaaac ctgggtctcaa 420
 cgccatcctg ggaccacag gtggaggcaa atcttcgtta ttagatgtct tagctgcaag 480

gaaagatcca agtggattat ctggagatgt tctgataaat ggagcaccgc gacctgccaa	540
tttcaaagt aattcagggt acgtggtaca agatgatggt gtgatgggca ctctgacggt	600
gagagaaaac ttacagttct cagcagctct tcggcttgca acaactatga cgaatcatga	660
aaaaaacgaa cggattaaca gggtcattca agagttaggt ctggataaag tggcagactc	720
caagggttga actcagttta tccgtgggtgt gtctggagga gaaagaaaaa ggactagtat	780
aggaatggag cttatcactg atccttccat cttgttcttg gatgagccta caactggctt	840
agactcaagc acagcaaagt ctgtcctttt gtcctgaaa aggatgtcta agcagggacg	900
aacaatcatc ttctccattc atcagcctcg atattccatc ttcaagttgt ttgatagcct	960
caccttattg gcctcaggaa gacttatggt ccacgggcct gctcaggagg ccttgggata	1020
ctttgaatca gctggttatc actgtgaggc ctataataac cctgcagact tcttcttgga	1080
catcattaat ggagattcca ctgctgtggc attaaacaga gaagaagact ttaaagccac	1140
agagatcata gagccttcca agcaggataa gccactcata gaaaaattag cggagattta	1200
tgtcaactec tccttctaca aagagacaaa agctgaatta catcaacttt ccgggggtga	1260
gaagaagaag aagatcacag tcttcaagga gatcagctac accacctcct tctgtcatca	1320
actcagatgg gtttccaagc gttcattcaa aaacttgctg ggtaatcccc aggctctat	1380
agctcagatc attgtcacag tcgtactggg actgggtata ggtgccattt actttgggct	1440
aaaaaatgat tctactggaa tccagaacag agctgggggt ctcttcttcc tgacgaccaa	1500
ccagtgtttc agcagtgttt cagccgtgga actctttgtg gtagagaaga agctcttcat	1560
acatgaatac atcagcggat actacagagt gtcactttat ttccttgga aactgttatc	1620
tgatttatta cccatgagga tgttaccaag tattatatatt acctgtatag tgtacttcat	1680
gttaggattg aaggcaaagg cagatgcctt cttcgttatg atgtttacc ttatgatggt	1740
ggcttattca gccagttcca tggcactggc catagcagca ggtcagagtg tggtttctgt	1800
agcaacactt ctcatgacca tctgttttgt gtttatgatg attttttcag gtctgttggt	1860
caatctcaca accattgcat cttggctgtc atggcttcag tacttcagca ttccacgata	1920
tggatttacg gctttgcagc ataatgaatt tttgggacaa aacttctgcc caggactcaa	1980
tgcaacagga aacaatcctt gtaactatgc aacatgtact ggcgaagaat atttggtaaa	2040
gcagggcatc gatctctcac cctggggcct gtggaagaat cacgtggcct tggcttgat	2100
gattgttatt ttctcacia ttgcctacct gaaattgtta tttcttaaaa aatattctta	2160
aatttcccct taattcagta tgatttatcc tcacataaaa aagaagcact ttgattgaag	2220
tattcaaaaa aaaaaaaaaa aaaaaaa	2247

<210> 27 <211> 655 <212> PRT <213> Homo sapien <400> 27

Met	Ser	Ser	Ser	Asn	Val	Glu	Val	Phe	Ile	Pro	Val	Ser	Gln	Gly	Asn
1				5					10					15	

Thr	Asn	Gly	Phe	Pro	Ala	Thr	Ala	Ser	Asn	Asp	Leu	Lys	Ala	Phe	Thr	20	25	30
Glu	Gly	Ala	Val	Leu	Ser	Phe	His	Asn	Ile	Cys	Tyr	Arg	Val	Lys	Leu	35	40	45
Lys	Ser	Gly	Phe	Leu	Pro	Cys	Arg	Lys	Pro	Val	Glu	Lys	Glu	Ile	Leu	50	55	60
Ser	Asn	Ile	Asn	Gly	Ile	Met	Lys	Pro	Gly	Leu	Asn	Ala	Ile	Leu	Gly	65	70	75
Pro	Thr	Gly	Gly	Gly	Lys	Ser	Ser	Leu	Leu	Asp	Val	Leu	Ala	Ala	Arg	85	90	95
Lys	Asp	Pro	Ser	Gly	Leu	Ser	Gly	Asp	Val	Leu	Ile	Asn	Gly	Ala	Pro	100	105	110
Arg	Pro	Ala	Asn	Phe	Lys	Cys	Asn	Ser	Gly	Tyr	Val	Val	Gln	Asp	Asp	115	120	125
Val	Val	Met	Gly	Thr	Leu	Thr	Val	Arg	Glu	Asn	Leu	Gln	Phe	Ser	Ala	130	135	140
Ala	Leu	Arg	Leu	Ala	Thr	Thr	Met	Thr	Asn	His	Glu	Lys	Asn	Glu	Arg	145	150	155
Ile	Asn	Arg	Val	Ile	Gln	Glu	Leu	Gly	Leu	Asp	Lys	Val	Ala	Asp	Ser	165	170	175
Lys	Val	Gly	Thr	Gln	Phe	Ile	Arg	Gly	Val	Ser	Gly	Gly	Glu	Arg	Lys	180	185	190
Arg	Thr	Ser	Ile	Gly	Met	Glu	Leu	Ile	Thr	Asp	Pro	Ser	Ile	Leu	Phe	195	200	205
Leu	Asp	Glu	Pro	Thr	Thr	Gly	Leu	Asp	Ser	Ser	Thr	Ala	Asn	Ala	Val	210	215	220
Leu	Leu	Leu	Leu	Lys	Arg	Met	Ser	Lys	Gln	Gly	Arg	Thr	Ile	Ile	Phe	225	230	235
Ser	Ile	His	Gln	Pro	Arg	Tyr	Ser	Ile	Phe	Lys	Leu	Phe	Asp	Ser	Leu	245	250	255
Thr	Leu	Leu	Ala	Ser	Gly	Arg	Leu	Met	Phe	His	Gly	Pro	Ala	Gln	Glu	260	265	270
Ala	Leu	Gly	Tyr	Phe	Glu	Ser	Ala	Gly	Tyr	His	Cys	Glu	Ala	Tyr	Asn	275	280	285
Asn	Pro	Ala	Asp	Phe	Phe	Leu	Asp	Ile	Ile	Asn	Gly	Asp	Ser	Thr	Ala	290	295	300
Val	Ala	Leu	Asn	Arg	Glu	Glu	Asp	Phe	Lys	Ala	Thr	Glu	Ile	Ile	Glu	305	310	315
Pro	Ser	Lys	Gln	Asp	Lys	Pro	Leu	Ile	Glu	Lys	Leu	Ala	Glu	Ile	Tyr	325	330	335
Val	Asn	Ser	Ser	Phe	Tyr	Lys	Glu	Thr	Lys	Ala	Glu	Leu	His	Gln	Leu	340	345	350
Ser	Gly	Gly	Glu	Lys	Lys	Lys	Lys	Ile	Thr	Val	Phe	Lys	Glu	Ile	Ser	355	360	365

Tyr	Thr	Thr	Ser	Phe	Cys	His	Gln	Leu	Arg	Trp	Val	Ser	Lys	Arg	Ser	370	375	380	
Phe	Lys	Asn	Leu	Leu	Gly	Asn	Pro	Gln	Ala	Ser	Ile	Ala	Gln	Ile	Ile	385	390	395	400
Val	Thr	Val	Val	Leu	Gly	Leu	Val	Ile	Gly	Ala	Ile	Tyr	Phe	Gly	Leu	405	410	415	
Lys	Asn	Asp	Ser	Thr	Gly	Ile	Gln	Asn	Arg	Ala	Gly	Val	Leu	Phe	Phe	420	425	430	
Leu	Thr	Thr	Asn	Gln	Cys	Phe	Ser	Ser	Val	Ser	Ala	Val	Glu	Leu	Phe	435	440	445	
Val	Val	Glu	Lys	Lys	Leu	Phe	Ile	His	Glu	Tyr	Ile	Ser	Gly	Tyr	Tyr	450	455	460	
Arg	Val	Ser	Ser	Tyr	Phe	Leu	Gly	Lys	Leu	Leu	Ser	Asp	Leu	Leu	Pro	465	470	475	480
Met	Arg	Met	Leu	Pro	Ser	Ile	Ile	Phe	Thr	Cys	Ile	Val	Tyr	Phe	Met	485	490	495	
Leu	Gly	Leu	Lys	Ala	Lys	Ala	Asp	Ala	Phe	Phe	Val	Met	Met	Phe	Thr	500	505	510	
Leu	Met	Met	Val	Ala	Tyr	Ser	Ala	Ser	Ser	Met	Ala	Leu	Ala	Ile	Ala	515	520	525	
Ala	Gly	Gln	Ser	Val	Val	Ser	Val	Ala	Thr	Leu	Leu	Met	Thr	Ile	Cys	530	535	540	
Phe	Val	Phe	Met	Met	Ile	Phe	Ser	Gly	Leu	Leu	Val	Asn	Leu	Thr	Thr	545	550	555	560
Ile	Ala	Ser	Trp	Leu	Ser	Trp	Leu	Gln	Tyr	Phe	Ser	Ile	Pro	Arg	Tyr	565	570	575	
Gly	Phe	Thr	Ala	Leu	Gln	His	Asn	Glu	Phe	Leu	Gly	Gln	Asn	Phe	Cys	580	585	590	
Pro	Gly	Leu	Asn	Ala	Thr	Gly	Asn	Asn	Pro	Cys	Asn	Tyr	Ala	Thr	Cys	595	600	605	
Thr	Gly	Glu	Glu	Tyr	Leu	Val	Lys	Gln	Gly	Ile	Asp	Leu	Ser	Pro	Trp	610	615	620	
Gly	Leu	Trp	Lys	Asn	His	Val	Ala	Leu	Ala	Cys	Met	Ile	Val	Ile	Phe	625	630	635	640
Leu	Thr	Ile	Ala	Tyr	Leu	Lys	Leu	Leu	Phe	Leu	Lys	Lys	Tyr	Ser		645	650	655	